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(54) **Modified proteins, modified interferons, alphas and beta, phosphorylated proteins and analogues, DNA sequences, applications and uses thereof**

Modifizierte Proteine, modifizierte Interferone alpha- und beta-phosphorylierte Proteine,
DNA-Sequenzen und Analoge und deren Verwendung

Protéines modifiées, interférons alpha et bêta modifiés, protéines phosphorylées et analogues,
séquences d'ADN, et leurs applications

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Description**FIELD OF THE INVENTION**

5 The present invention relates to the field of recombinant DNA technology and to means and methods of utilizing this technology to synthesize functional proteins or polypeptides useful in biomedical, medical, biochemical applications including diagnostics, prophylactics and therapeutics.

BACKGROUND OF THE INVENTION

10 Radio-labelled proteins have numerous medical, biological, clinical, scientific and other applications. Interferons, specifically, labelled with ¹²⁵I have been used for binding and crosslinking studies (1, 17, 27-32, 34-37).¹ Human IFN- α 's, - β , and -gamma have all been radio-iodinated by various procedures (reviewed in Pestka et al (2)). However, proteins labelled with radioactive iodine have serious well-known disadvantages and hazards.

15 The study of cell surface receptors for the interferons requires radio-labelled interferons, such as interferons labelled with ¹²⁵I, with high biological and high radio-specific activity. Several years ago, it was found that interferon gamma² can be phosphorylated to very high radio-specific activity while retaining biological activity (3, 4). Thus, [³²P]Hu- and Mu-IFN-gamma were used for studying the human and murine IFN-gamma receptors, respectively (5, 6, 9). These studies were carried out by phosphorylating human and murine interferon gamma (Hu- and Mu-IFN-gamma) with cyclic
20 AMP-dependent protein kinase from bovine heart muscle and [gamma-³²P]ATP (3). These phosphorylated and ³²P-labelled interferons have provided valuable reagents (3, 4) of high radio-specificity to study cell surface receptors (5, 6) and to identify the chromosome encoding the gene for Hu-IFN-gamma (7, 8) and Mu-IFN-gamma (9) receptors. For all of these studies and applications, interferons which are phosphorylated are most useful. Several reports identified the phosphorylation sites of Hu- and Mu-IFN-gamma as serine residues near the COOH termini (4, 5, 10, 11).

25 However, under conditions used for the phosphorylation of IFN-gamma, it was reported that Hu-IFN- α A and Hu-IFN- β cannot be phosphorylated by the cyclic AMP (cAMP)-dependent protein kinase (2, 3). A review of the phosphorylation of the various classes or groups of interferons and other proteins (1, 3, 4, 20, 21, 22, 38, 39, 40, 64) confirms that researchers have not successfully phosphorylated Hu-IFN- α or Hu-IFN- β under conditions under which gamma interferons have been phosphorylated. It has been reported indeed that recombinant IFN- α and IFN- β were not phos-
30 phorylated (3) and as a consequence it was uncertain whether an available site was present.

In the light of problems with iodinated compounds and limitations for use of iodinated IFN-gamma, it is understandable that there is a keen interest and need in making available phosphorylated Hu-IFN- α and - β which can be labelled for numerous practical, scientific and commercial applications.

35 Likewise, there is such interest and need for other phosphorylated -- and labelled -- polypeptides which are not available yet in such chemical configurations. For example, a phosphorylatable tumor necrosis factor (TNF) would be valuable to study the receptor for TNF. TNF is not phosphorylatable with the cAMP-dependent bovine heart kinase. Indeed, it has been reported that interest in protein phosphorylation has increased enormously over the past few years (38, 39).

The invention as will be described in detail hereinafter contributes to meeting these and other needs.

40 By way of further background to the invention, the term "interferon" describes a family of animal proteins which possess antiviral, antiproliferative and other potentially useful properties. There appear to be three major classes of interferons: leukocyte (or alpha interferon), fibroblast (or beta interferon) and immune (or gamma interferon) (1, 2). Detailed description of interferons is found in various publications including in references 1, 2, U.S. Patent Nos. 4,727,138; 4,734,491; 4,737,462, and many others; various hybrid human leukocyte interferons are described in U.S.
45 Patent No. 4,414,150 and in reference 46. In general the standard class of human IFN- α 's are polypeptides of 165-166 amino acids (see reference 1 for details of human and non-human interferon- α species); some species have been isolated that lack the 10 COOH-terminal amino acid residues; and some species of IFN- α are glycosylated. The amino acid sequences of Hu-IFN- α species and of Hu-IFN- β derived from cDNA or genomic DNA sequences are described in (1, Section I). Recombinant DNA-derived interferons including Hu-IFN- α , - β , and -gamma and corresponding inter-
50 ferons from other animal species are likewise well described (1, 2). Various modifications of human and murine interferons have been reported. New non-natural human and murine interferons with often markedly changed biological properties have been constructed (1, 24, 45). The terminology "non-natural" is a term of art which refers to recombinant DNA interferons obtained by altering the nucleotide sequence of coding cDNAs (45).

¹ The scientific publications, patents or other literature ("publications") to which reference is made herein are referenced by numerals and identified further towards the end of this text. All of these publications are incorporated herein by reference.

² The abbreviations used have followed standard nomenclature as described in detail in Methods of Enzymology, Interferons, Vol. 119, Part C, Edited by Sidney Pestka, Section I, Introduction (Reference 25). In brief, interferon alpha, beta, and gamma are designated IFN- α , IFN- β , and -gamma, respectively. The species of origin is designated by a prefix Hu, Mu, Bo, etc. for human, murine, or bovine species, respectively, as Hu-IFN- α , Hu-IFN- β , or Hu-IFN-gamma, for example.

The term "Hu-IFN- α " as used herein is intended to include all different species of alpha interferons. A large number of DNA sequences corresponding to the interferons from various species have been isolated and identified. Likewise various IFN- β s and IFN-gamma(s) are disclosed. The invention encompasses all of these members of the family (reference 1, pages 5-14).

The term "native" as used herein refers to the proteins, e.g., interferons, which proteins are naturally produced; "synthetic" and "non-natural" refers to proteins produced by synthetic or DNA-recombinant procedures, either type which do not contain a phosphorylatable site (or where the phosphorylatable site is inaccessible, for instance due to the configuration of the protein), which protein in accordance with the invention is to be phosphorylated.

This invention contemplates and includes all interferons native, natural, modified, or recombinant DNA interferon-like proteins which are modifiable by introduction of one or more phosphate or analog groups. All of these interferons and others known in the art or to be known are within the contemplation of the invention. The present invention is principally concerned with various modified proteins or polypeptides, and alpha and beta interferons.

When reference is made to IFN-alpha, the term is intended to cover and include the various alpha species.

The term "modified" is used in this invention broadly, and means for instance, when reference is made to proteins, a protein which has been provided with a phosphorylatable site or provided with a phosphorus label (or analog label). The nucleotide sequences which code for such amino acid sequences which contain a putative phosphorylation site are also designated as "modified", when appropriate.

The term "unphosphorylatable" protein means a protein which normally has not been phosphorylatable (or phosphorylated) for whatever reason, e.g., either because it does not contain a putative phosphorylatable site and correspondingly, the DNA sequence which codes for the protein does not contain the DNA sequence coding for the putative amino acid recognition sequence; or because such site is not accessible for phosphorylation.

The term "provided with" or "having provision(s) for" or like terminology is used in this invention broadly, and means both "fused" and "inserted". Illustrative are the hybrid-fused Hu-IFN- α /gamma (illustrated in FIG. 1) and Hu-IFN- α -P1, -P2 and -P3 (illustrated in FIG. 8), respectively. Thus, the nucleotide insert can be within the coding region of the gene at one end thereof or anywhere within the coding region. These variants are all considered to be within the term "modified," which can refer to the amino acid sequence or to the nucleotide sequences, as will become apparent from the description hereinafter.

The term "comprises" or "comprising" covers and includes all situations regardless where the amino acid recognition sequence (or the nucleotide sequence coding for it) is located.

By way of further background in the method of the invention, phosphorylation is carried out by means of a protein kinase. Protein kinases catalyze the transfer of the gamma phosphoryl group of ATP to an acceptor protein substrate. However, as described herein the invention is not limited to kinases for which the acceptor site is a particular amino acid (like serine) but includes also those for which the site is another amino acid in the sequence, and in general includes protein kinases as a whole.

The term "protein" (or polypeptide) as used herein is intended to include glycoproteins (as well as proteins having other additions). A case in point is that of natural Hu-IFN- β which has been shown to be a glycoprotein; when produced in *E. coli* by recombinant DNA techniques, Hu-IFN- β is not glycosylated. Glycosylated interferons have been reported to be obtained by expressing the proteins in animal cells or in yeast (as is discussed in reference 1 at pps. 383-433 and 453-464; and in references 48-55, 84-92).

The term "biological activities" or like terms as used herein in conjunction with proteins is intended to be interpreted broadly. In the case of the interferon-like proteins, it includes all known (or to be discovered) properties including properties specific to Hu-IFN- α 's or to Hu-IFN- β or common to both, such as their antiviral activity and their capability to modulate antigens of the major histocompatibility complex (MHC); in particular to induce an increase in surface expression of class I MHC antigens, including β_2 -microglobulin.

"Functional" proteins are proteins which have a biological or other activity or use.

The term "active areas" or "biologically active" areas or segments or equivalent terminology often refers to the presence of a particular conformation or folding of the protein molecule, or for instance, to specific disulfide bridges between specific amino acids in the sequence, but of course is not limited thereto.

The term "vector" as used herein means a plasmid, a phage DNA, or other DNA sequence that (1) is able to replicate in a host cell, (2) is able to transform a host cell, and (3) contains a marker suitable for identifying transformed cells.

Throughout the description of the invention and the claims, and following convention, the "singular" includes the "plural"; for instance, a phosphorylatable or phosphorylation site, means at least one such site, unless indicated otherwise.

Other terminology used herein will become apparent from the description which follows.

BRIEF DESCRIPTION OF THE PRIOR ART

Background references for the subject invention are referred to within the body and towards the end of the text.

As representative of United States patents which relate to interferon, the following may be mentioned:

U. S. Patent No. 4,503,035 to Pestka et al relates to human leukocyte interferon as a homogeneous protein species, such as species α_2 , α_2 , and β_1 , and others. For a discussion of terminology of natural and recombinant interferons see references 1 (pps. 3-23), 24, 102, and 103 (footnote p. 112 and text);

It is noteworthy that the above reviewed patent literature does not address or disclose human intetferons which have phosphorylated groups (or isotopes thereof).

SUMMARY AND GENERAL CONCEPTS OF THE INVENTION

In a broad sense, the invention contemplates labellable and labelled proteins, e.g. radio-labellable and radio-labelled proteins, and DNA and cDNA molecules encoding the radio-labellable proteins.

The invention encompasses recombinant DNA sequences which encode functional proteins having one or more putative phosphorylation sites; expression vectors for expressing the functional protein; transformed host, methods of expressing the modified proteins and the modified proteins.

In one embodiment, the invention provides radioactive-labelled human interferons and labelled proteins; phosphorylatable modified Hu-IFN- α (Hu-IFN- α -P) which can be phosphorylated to high radio-specific activity with retention of biological activity; other human interferons modified with various isotopes of phosphorus (e.g., ^{32}P , ^{33}P), or with sulfur (e.g., ^{35}S , ^{38}S); labelled proteins with phosphorus or analogs. In accordance with the invention, the human interferons and modified proteins may have single or multiple radioactive labels.

The invention also provides such interferons and proteins made by recombinant DNA techniques, including the Hu-IFN- α -P human interferons radio-labelled with phosphorus or with sulfur, and recombinant DNA-produced radio-labelled polypeptides and proteins.

The invention further provides DNA sequences encoding a functional protein which possesses one or more labelling sites and is sufficiently duplicative of human interferons for the protein sequences to possess at least one of the biological properties of interferons (like antiviral, cell growth inhibiting, and immunomodulatory properties). Further, there is provided a recombinant-DNA containing a coding sequence for a putative recognition site for a kinase; the recombinant expression vector; the host organisms transformed with the expression vector that includes the DNA sequence and an expressed modified protein. In the invention, there is used a method involving site-specific mutagenesis for constructing the appropriate expression vector, a host transformed with the vector and expressing the modified proteins, in particular the modified human interferons.

The invention provides in one of its several embodiments DNA sequences which encode one or more putative phosphorylation sites, which sequences encode functional proteins each of which possesses at least one putative phosphorylation site and each of which possesses at least one of the biological properties of Hu-IFN- α or - β ; also expression vectors for expression of the functional modified Hu-IFN- α or - β under the control of a suitable promoter such as the lambda P_c promoter or others described hereinafter; also the biologically active phosphorylated Hu-IFN- α and - β .

Several interesting and useful applications of these modified human interferons and proteins are also disclosed by the description.

The invention also contemplates interferons or proteins other than the Hu-IFN- α or - β , which are modified by addition of phosphorylation sites which allow for and are labelled to higher radio-specific activities than the corresponding interferons with a single phosphorylation site. By "addition" of phosphorylation sites, there is also intended in accordance with the invention, to include interferons or proteins in which a phosphorylation site heretofore unavailable or inaccessible, has been modified to make the phosphorylation site available.

The invention further contemplates interferons, especially Hu-IFN- α , phosphorylated by appropriate kinases on amino acid residues other than on the serine residue, like on threonine and/or tyrosine residues, and the DNA sequences which code for one or more putative phosphorylation sites, which sequences code for these interferons.

In accordance with the invention, it is sufficient that a portion of the phosphorylation recognition sequence, as opposed to the entire sequence, be added when the natural protein sequence contains the remaining (or other complementary) amino acids of said recognition sequence (e.g., Arg-Arg-Ala-Ser). In such embodiment of the invention, from 1 through 4 amino acids of the sequence (in the case of Arg-Arg-Ala-Ser-Val) can be supplied to the protein, thereby constituting the complete, necessary and Ser-containing recognition sequence. An illustration can be observed in a comparison between species Hu-IFN- α -P1 and -P2 (in FIG. 8), wherein the natural interferon sequence contributes one Arg to the phosphorylation recognition sequence in Hu-IFN- α -P2 when constructed in accordance with the invention.

In Hu-IFN- α -P3, a coding sequence (and thus an additional amino acid sequence) has been supplied with the nucleotide sequence coding for the recognition sequence positioned downstream of the natural sequence coding for Hu-IFN- α . Thus, Hu-IFN- α -P3 is an illustration where an additional amino acid sequence is positioned between the recognition sequence and the natural amino acid sequence of Hu-IFN- α .

This illustrates the versatility of the invention for positioning the nucleotide sequence which encodes the amino acid

recognition sequence containing a putative phosphorylation site.

Thus, in accordance with the invention, there is constructed a nucleotide sequence that codes for the necessary number and specific amino acids required for creating the putative phosphorylation site.

From the above observation, the same principles are applicable to construct any amino acid sequences other than the particular amino acid recognition sequence illustrated above.

In the situations where the phosphorylation site is other than serine (as illustrated above), the DNA sequence codes for part or all of the appropriate amino acid sequence containing the putative recognition site containing threonine, tyrosine, etc. Thus, where in any particular protein one or more amino acids (at any position of the amino acid sequence) are the same as that of an amino acid recognition sequence for a kinase, it is sufficient to add (or modify) those complementary amino acids of the amino acid recognition sequence to complete that sequence. This is accomplished by constructing a DNA sequence which codes for the desired amino acid sequence. There may indeed be situations where such addition (or modification) is a more desirable procedure as where it is important to retain the integrity of the protein molecule to be modified (for instance, to minimize risks of affecting a particular activity, e.g., biological), or for simplicity of the genetic manipulations, or because either or both termini or other positions are more accessible.

The kinase recognition sequence may be positioned at either termini or other position of the DNA coding sequence, irrespective of the specific phosphorylated amino acid.

In accordance with the invention, phosphorylation of the phosphorylatable site of the protein can be performed by any suitable phosphorylation means. Phosphorylation and dephosphorylation of proteins catalyzed by protein kinases and protein phosphatases is known to affect a vast array of proteins (21). A large number of protein kinases have been described (20, 21, 22, 38, 39, 47, 64, 100, 101, 108-112) and are available to one skilled in the art for use in the invention. Such protein kinases may be divided into two major groups: those that catalyze the phosphorylation of serine and/or threonine residues in proteins and peptides and those that catalyze the phosphorylation of tyrosine residues (see 21, 22, 38, 64, 108, for example). These two major categories can be subdivided into additional groups. For example, the serine/threonine protein kinases can be subdivided into cyclic AMP (cAMP)-dependent protein kinases, cyclic GMP (cGMP)-dependent kinases, and cyclic nucleotide-independent protein kinases. The recognition sites for many of the protein kinases have been deduced (21, 22, 38, 64, 111 present illustrative examples).

In short synthetic peptides CAMP-dependent protein kinase recognize the sequence Arg-Arg-Xxx-Ser-Xxx, where Xxx represents an amino acid (21). As noted above, the CAMP-dependent protein kinase recognizes the amino acid sequence Arg-Arg-Xxx-Ser-Xxx (21), but also can recognize some other specific sequences such as Arg-Thr-Lys-Arg-Ser-Gly-Ser-Val (111). Many other protein serine/threonine kinases have been reported (21, 100, 101, 108-112) such as glycogen synthase kinase, phosphorylase kinase, casein kinases I and II, pyruvate dehydrogenase kinase, protein kinase C, and myosin light chain kinase.

Protein kinases which phosphorylate and exhibit specificity for tyrosine (rather than for serine, threonine, or hydroxyproline) in peptide substrates are the protein tyrosine kinases (PTK). Such PTKs are described in the literature (22, 64). The PTKs are another class of kinases available for use in the invention.

Another available class of kinases are the cyclic GMP-dependent (cGMP-dependent) protein kinases. The cGMP-dependent protein kinases exhibit substrate specificity similar to, but not identical to the specificity exhibited by CAMP-dependent protein kinases. The peptide Arg-Lys-Arg-Ser-Arg-Lys-Glu was phosphorylated at serine by the cGMP-dependent protein kinase better than by the cAMP-dependent protein kinase (21, 22, 113). It has also been shown that the cAMP-dependent protein kinase can phosphorylate hydroxyproline in the synthetic peptide Leu-Arg-Arg-Ala-Hyp-Leu-Gly (114).

Casein kinases, widely distributed among eukaryotic organisms and preferentially utilizing acidic proteins such as casein as substrates, have been classified into two groups, casein kinases I and II (21). Casein kinase II phosphorylated the synthetic peptide Ser-Glu-Glu-Glu-Glu-Glu (115). Evaluation of results with synthetic peptides and natural protein substrates revealed that a relatively short sequence of amino acids surrounding the phosphate acceptor site provides the basis for the specificity of casein kinase II (118). Accordingly, the acidic residues at positions 3 and 5 to the carboxyl-terminal side of the serine seem to be the most important. Serine was preferentially phosphorylated compared to threonine. In another study (117), the peptide Arg-Arg-Arg-Glu-Glu-Glu-Thr-Glu-Glu-Glu was found to be a specific substrate for casein kinase II; however, Arg-Arg-Arg-Glu-Glu-Glu-Ser-Glu-Glu-Glu was a better substrate (118); and Arg-Arg-Arg-Asp-Asp-Asp-Ser-Asp-Asp-Asp was a better substrate than Arg-Arg-Arg-Glu-Glu-Glu-Ser-Glu-Glu-Glu. Thus, aspartate is preferred over glutamate (118). Acidic residues on the COOH-terminal side of the serine (threonine) are as far as known today absolutely required; acidic residues on the amino-terminal side of the serine (threonine) enhance phosphorylation, but are not absolutely required: thus, Ala-Ala-Ala-Ala-Ala-Ala-Ser(Thr)-Glu-Glu-Glu served as a substrate for casein kinase II, but was less effective than Ala-Ala-Ala-Glu-Glu-Glu-Ser(Thr)-Glu-Glu-Glu (118) (the designation Ser(Thr) means serine or threonine). Casein kinases I and II phosphorylate many of the same substrates (21) although casein kinase I did not phosphorylate any of the decamer peptide substrates noted here (118). It was concluded from studies with a variety of synthetic peptides that the sequence Ser-Xxx-Xxx-Glu (and by inference Ser-Xxx-Xxx-Asp) may represent one class of sequences that fulfill the minimal requirements for recognition by casein

kinase II although some other peptides and sequences may also suffice (see 118 for a detailed discussion).

As noted above, other kinases have been described. The mitogen-activated S6 kinase phosphorylates the synthetic peptide Arg-Arg-Leu-Ser-Ser-Leu-Arg-Ala (109) as does a protease-activated kinase from liver (21, 109). The rhodopsin kinase catalyzes the phosphorylation of the peptide Thr-Glu-Thr-Ser-Gln-Val-Ala-Pro-Ala (21). Other protein serine/threonine kinases have been described and their sites of phosphorylation elucidated (21).

The substrate specificity of tyrosine kinases have also been reported (64, pages 920-921; 110). A variety of synthetic or natural peptide substrates have been described (64, 110).

Thus, one skilled in the art has quite an adequate selection of available kinases for use in the invention, which have relatively high specificity with respect to the recognition process, but some flexibility to the specific sequence of the amino acid recognition site. Such kinases provide means for phosphorylation of putative phosphorylation sites in the desired proteins.

The selection of the position of the molecule best suited for the modification depends on the particular protein (and its configuration). Where multiple putative phosphorylation sites (and phosphorylatable sites) are to be included in the modified protein, one would consider the potential availability of either or both ends and other positions of the molecule for providing the amino acid recognition sequence. Thus, in accordance with the invention, phosphorylation recognition sequences can be introduced at any point in a naturally occurring protein sequence providing such introduced sequences do not adversely affect biological activity where such activity is desired.

Once the recognition site for a particular protein kinase is identified, the invention provides a method for making by recombinant-DNA techniques the DNA sequence which encodes the recognition site for that kinase within, fused or linked to the DNA sequence encoding the functional protein which is to contain the corresponding putative labelling site.

The invention contemplates and includes any protein which is radio-labellable by the methods of this invention and which possesses at least one of the properties of the corresponding unlabelled (or unlabellable) protein. In accordance with the invention, the non-phosphorylated (or non-phosphorylatable) protein is modified to introduce into the amino acid sequence the putative phosphorylatable site; this is performed after having modified the DNA sequence encoding the amino acid sequence of the protein with the DNA sequence (part or all) which codes for the putative phosphorylated site. In the case of interferons, the invention includes all interferons natural or "non-natural" interferons, including such structurally modified interferon species which have been reported in the literature (such as hybrid interferons, modified interferons) as discussed above, and other modified interferons which will be reported in the future.

Natural and "non-natural" (including modified) interferon species have a variety of biological activities and such activities are known to occur in different ratios; thus, the invention contemplates not only radio-labelled interferons which have any one of these properties (and in any ratio), but also biological or other properties not yet identified in the known interferon species. It is recognized that the phosphorylation may modify one or more of the properties of the protein to one degree or another (see 47, 100, 101, for example). Indeed there are situations where the properties may be enhanced or developed where they were not detectable prior to modification of the protein.

The invention also provides particularly interesting labellable and labelled proteins like phosphorylated antibodies (especially monoclonal antibodies, hybrid antibodies, chimeric antibodies or modified antibodies), hormones, and "modified" streptavidin. The modified streptavidin can be bound to individual biotinylated antibodies, each streptavidin being modified by single or multiple phosphorylated groups, which product has greatly enhanced radiation and therefore diagnostic and therapeutic potential.

The invention also provides a hybrid interferon protein Hu-IFN- α A/gamma constituted of Hu-IFN- α A to which there is fused the COOH-terminal 16 amino acid region of Hu-IFN-gamma, which contains a putative phosphorylation site, and the hybrid interferon fusion protein labelled with phosphorus. The fusion protein was synthesized with an expression vector constructed by oligonucleotide-directed mutagenesis. The invention also provides the DNA coding sequence for the fused hybrid interferon protein, expression vectors and the transformed microorganisms, e.g. *E. coli* host and other suitable hosts described below.

The foregoing is not intended to have identified all of the aspects or embodiments of the invention nor in any way to limit the invention. The invention is more fully described below.

The accompanying drawings, which are incorporated and constitute part of the specification, illustrate embodiments of the invention, and together with the description, serve to explain the principles of the invention.

BRIEF DESCRIPTION OF THE DRAWINGS

The Figures first illustrate the fused-hybrid interferon (FIGS. 1-7) and then the other modified interferons made in accordance with the invention (FIGS. 8-13).

FIG. 1 is a schematic illustration of a hybrid interferon Hu-IFN- α A/gamma of Hu-IFN- α A and Hu-IFN-gamma, the Hu-IFN-gamma segment representing the 16 COOH-terminal amino acids and containing the putative phosphorylation site.

FIG. 2 shows an outline of the construction of the fusion protein from Hu-IFN- α A and Hu-IFN-gamma, the

Hu-IFN- α /gamma.

The EcoRI fragment from pRC23/IFN-gamma was ligated into the BstEII site of pXZ-6 (pIFN- α A) by blunt-end ligation to yield pXZ-7. The EcoRI fragment of pXZ-7 was inserted into M13mp18 to yield M13mp18/IFN- α A/IFN-gamma. Oligonucleotide-directed deletion was performed on this latter construct with the oligonucleotide shown to yield a re-

combinant containing the proper deletion (M13mp18/IFN- α A/gamma). The EcoRI fragment from this M13 recombinant

replaced the EcoRI insert of pXZ-6 to yield pXZ-8 (pIFN- α A/gamma), the expression plasmid encoding the Hu-IFN- α /gamma fusion protein. The proper construction was confirmed by sequencing the recombinant. T4-t represents the phage T4 transcription terminator (56, 57); Ap, ampicillin; cip, calf intestinal phosphatase.

FIG. 3 shows the nucleotide sequence encoding and amino acid sequence of the hybrid Hu-IFN- α /gamma.

FIG. 4 shows a polyacrylamide gel electrophoresis of the purified proteins. Lane 1 represents Hu-IFN- α A; lane 2, the fused protein Hu-IFN- α /gamma; and lane 3, Hu-IFN-gamma. The molecular weight standards (STD) are shown in the right column. The gels were developed with a silver stain.

FIG. 5 shows a polyacrylamide gel electrophoresis of phosphorylated Hu-IFN-gamma and the fusion protein

Hu-IFN- α /gamma. The left panel of the Figure represents the gel stained with Coomassie blue. The right panel represents the autoradiograph of the same gel as shown on the left panel. The heavy stained bands at MW_R 69,000 represent bovine serum albumin which was added to the phosphorylation reactions.

FIG. 6 shows the binding of [³²P]Hu-IFN- α /gamma to Daudi cells. The binding of [³²P]Hu-IFN- α /gamma to Daudi cells was performed as described under "Experimental Procedures". Specific binding (●) represents the difference between total binding (○) and non-specific binding (Δ). Non-specific binding represents the binding in the presence of excess unlabelled Hu-IFN- α A. The specific activity of [³²P]Hu-IFN- α /gamma was about 141 μ Ci/ μ g.

FIG. 7 shows the binding of [³²P]Hu-IFN- α /gamma to MDBK cells. The symbols and explanations are the same as in FIG. 6.

FIG. 8 shows nucleotide and amino acid sequences of the COOH terminus of Hu-IFN- α A, Hu-IFN- α A-P1, -P2, and -P3. The phosphorylation sites recognized by the cAMP-dependent protein kinase created in Hu-IFN- α A-P1, -P2, and -P3 are shown in rectangles. The sites contain the recognition sequence Arg-Arg-Ala-Ser-Leu or Arg-Arg-Ala-Ser-Val (for phosphorylation of this Ser) for the cAMP-dependent bovine heart kinase together with additional amino acids. The nucleotide sequences and amino acid sequences of the interferons are aligned for comparison.

FIG. 9 shows the construction of expression plasmids for Hu-IFN- α A containing phosphorylation sites.

FIG. 10 shows an analysis of Hu-IFN- α A-P proteins by SDS-polyacrylamide gel electrophoresis. The gel was stained with silver. Molecular weight standards are shown on the left most lane.

FIG. 11 shows an autoradiograph of SDS-polyacrylamide gel electrophoresis of the phosphorylated proteins. Hu-IFN-gamma serves as an internal molecular weight standard as in FIG. 10.

FIG. 12 shows the binding of [³²P]Hu-IFN- α A-P proteins to cells. The inset represents Scatchard analysis of the specific binding data. "B" represents the radioactivity of ligands bound to cells and "F" is the radioactivity of free or unbound ligands. (A) and (B): binding [³²P]Hu-IFN- α A-P1 to Daudi and MDBK cells, respectively; (C) and (D): binding of [³²P]Hu-IFN- α A-P2 to Daudi and MDBK cells, respectively.

FIG. 13 shows the electrophoretic analysis of crosslinked [³²P]Hu-IFN- α A-P1, -P2 and -P3 receptor complexes.

The Figures are explained in greater detail hereinbelow.

DESCRIPTION OF SPECIFIC EMBODIMENTS

In accordance with the invention, proteins which are normally not phosphorylatable can be modified to render them phosphorylatable. In particular, it was of great interest in conjunction with this invention to determine whether and how one could achieve the phosphorylation of a particular protein, the interferon Hu-IFN- α A, which as opposed to Hu-IFN-gamma could not be phosphorylated (as discussed above). The methodology to achieve this result (especially without loss of its biological activity) has provided the potential to modify other proteins and render them phosphorylatable. Thus, in accordance with the invention, Hu-IFN- α A has been used as a model for phosphorylation of proteins in general.

Hu-IFN- α A has been modified in accordance with the invention to yield two embodiments, which will be described hereinafter. Both embodiments are applicable to proteins in general and one skilled in the art will select that most suitable under the particular circumstances.

The first embodiment is the construction of a phosphorylatable hybrid or fused protein, Hu-IFN- α /gamma; the second embodiment is a phosphorylatable modified Hu-IFN- α A with an insert more fully described below.

The following describes illustrative, but not limiting, specific embodiments.

Hu-IFN- α /gamma, a hybrid protein of Hu-IFN- α A and the COOH-terminal 16 amino acids of Hu-IFN-gamma.

The following describes a hybrid or fused protein of Hu-IFN- α A and Hu-IFN-gamma. The fused-hybrid protein was

expressed by an expression vector constructed by oligonucleotide-directed mutagenesis. The construction of the expression vector, the expression and purification of the protein, the phosphorylation of the functional hybrid protein and the binding of [32 P]Hu-IFN- α /gamma to cells (MDBK) are described below. The hybrid protein has antiviral activity.

The fusion protein prepared in accordance with the invention comprises Hu-IFN- α A to which the COOH-terminal 16 amino acids of Hu-IFN-gamma was fused. The protein was prepared by constructing an expression vector by oligonucleotide-directed mutagenesis. The hybrid protein Hu-IFN- α A/gamma was expressed under the control of the phage lambda P_L promoter. The protein was purified with a monoclonal antibody against Hu-IFN- α or the COOH-terminal amino acid sequence of Hu-IFN-gamma. The purified protein exhibited a single major band on sodium dodecyl sulfate-polyacrylamide gel electrophoresis and has antiviral activity on human and bovine cells. Unlike Hu-IFN- α A, but similar to Hu-IFN-gamma, it was found that the hybrid Hu-IFN- α A/gamma can be phosphorylated by [gamma- 32 P]ATP and cAMP-dependent protein kinase. The phosphorylated molecule binds to the IFN- α / β receptor.

The introduction of a phosphorylation site into Hu-IFN- α A by fusion of the region of Hu-IFN-gamma which contains the putative phosphorylation site provides a new reagent for studies of receptor binding, pharmacokinetics and other studies where labelled interferons are useful.

Thus, in accordance with the invention, a hybrid protein was constructed by fusing the COOH-terminal end of Hu-IFN-gamma to the COOH terminus of Hu-IFN- α A to yield a hybrid protein which contains the putative phosphorylation site. The phosphorylated protein was found to be biologically active.

The construction of a plasmid for expression of hybrid Hu-IFN- α A/gamma was carried out as follows.

As shown in FIG. 2, the DNA sequences for Hu-IFN- α A and Hu-IFN-gamma were cloned into M13mp18. Plasmid pXZ-8 encoding a fusion protein was constructed by oligonucleotide-directed deletion as shown.

Plasmid pXZ-8 is deposited with the ATCC under Accession number 40510, and designated as pHu-IFN- α A/gamma.

To screen for the plasmid recombinant with the proper deletion, colony hybridization was performed on 200 colonies with the oligomer as probe (58) with the observation that 53 colonies were positive. Of the 53 positive colonies, 36 were analyzed by restriction digestion with *Eco*RI with the result that only two colonies contained the expected restriction fragments. DNA sequencing confirmed that the 5' end of the sequence coding for the 16 COOH-terminal amino acids of Hu-IFN-gamma was fused to the 3' end of the coding sequence for Hu-IFN- α A (FIGS. 1, 2, and 3). A 732 bp fragment was deleted in the correct recombinant. The gene for hybrid Hu-IFN- α A/gamma was constructed and expressed under control of the lambda P_L promoter. The sequence of hybrid Hu-IFN- α A/gamma consists of 181 amino acids (FIG. 3).

As discussed further below, other promoters in various expression vectors can also be used.

The characterization and phosphorylation of hybrid Hu-IFN- α A/gamma was performed as follows.

Hu-IFN- α A/gamma was induced when the *E. coli* RR1(pRk248c|ts857) cells or *E. coli* AR68(pRK248c|ts857) cells harboring the plasmid pXZ-8 were grown at 42°C and the hybrid protein was purified by monoclonal antibody affinity chromatography as described hereinafter. The purified protein exhibited a specific activity of 4×10^8 units/mg. On SDS-polyacrylamide gel electrophoresis in the presence of 2-mercaptoethanol, it exhibited a major single band which migrated slower than Hu-IFN- α A and almost at the same position of Hu-IFN- α A-P3, another Hu-IFN- α A derivative with 182 amino acids as shown in FIGS. 4 and 5. It was observed that Hu-IFN- α A/gamma is not as stable as Hu-IFN- α A. During experimental procedures, purification steps, and storage, degradation of the protein extracted from *E. coli* RR1 was observed; much less degradation was observed when *E. coli* AR68 was used as the host cell.

It was found that hybrid Hu-IFN- α A/gamma can be phosphorylated with [gamma- 32 P]ATP and bovine heart muscle cAMP-dependent protein kinase. The specific radioactivity of [32 P]Hu-IFN- α A/gamma was about 5 μ Ci/ μ g protein in preparations isolated from *E. coli* RR1; in preparations of Hu-IFN- α A/gamma isolated from *E. coli* AR68, the specific radioactivity was increased to about 141 μ Ci/ μ g, presumably reflecting greater integrity of the carboxyl terminus. These values are similar to the range of values obtained for [125 I]IFN- α A (59-62). [32 P]IFN- α A/gamma migrates as a single species at the same position as unlabelled IFN- α /gamma on SDS-polyacrylamide gel electrophoresis in the presence of 2-mercaptoethanol (FIG. 5).

The binding of [32]Hu-IFN- α A/gamma to cells was as follows.

Bovine MDBK cells were used for binding studies because Hu-IFN- α A exhibits high antiviral activity on this cell line and binds to it (16, 28, 63), whereas Hu-IFN-gamma does not. To determine whether Hu-IFN- α A/gamma binds specifically to bovine MDBK and human Daudi cells, cells were incubated with various concentrations of [32 P]Hu-IFN- α A/gamma in the absence and presence of unlabelled Hu-IFN- α A. Specific binding of [32 P]Hu-IFN- α A/gamma to both MDBK and Daudi cells is shown in FIGS. 6 and 7.

These results with [32 P]Hu-IFN- α A/gamma indicate that the fusion of the phosphorylation site of Hu-IFN-gamma to the COOH terminus of Hu-IFN- α A, which itself cannot be phosphorylated, did not destroy the capability of the protein to bind. The Hu-IFN- α A/gamma protein also retained its activity.

This description of the first embodiment of the invention shows the successful construction of a hybrid DNA sequence capable of coding for an amino acid sequence containing a putative phosphorylation site, the construction of a suitable expression vehicle and of a transformed host which readily expressed the hybrid protein with an intact putative phosphorylation site, and its phosphorylation.

The second principal embodiment of the invention reached the initial objective in a somewhat different manner; yet it still resulted in a modified interferon which is phosphorylatable.

While in the first above-described embodiment, the DNA sequence coding for the entire protein which is not phosphorylatable, in this case Hu-IFN- α A, is modified by fusion at the end of its sequence to a sequence which codes for a putative phosphorylatable amino acid sequence, the second embodiment does not require the fusion to be at the end of the sequence and, in fact, does not require a fusion step.

Thus, the second embodiment shows the wider applicability of the invention to proteins in general.

In the second embodiment the resulting protein has been modified by insertion of an appropriate phosphorylatable sequence.

The second embodiment will be described hereinafter.

Hu-IFN- α Having Inserted Labellable Sites.

An important concept of the invention is the incorporation of a kinase-recognizable amino acid sequence into selected proteins (or converting an amino acid sequence which is not recognizable by a protein kinase into a sequence which is so recognized), and labelling the proteins which contain the putative kinase-recognizable phosphorylation site by attaching a selected radioactive label by means of the catalytic action of the kinase.

Generally, these steps will be performed at the level of the nucleotide sequence by DNA recombinant procedures. After the amino acid consensus sequence recognizable by a protein kinase is identified, the nucleotide sequence encoding this amino acid sequence can be defined and constructed. For purposes of this disclosure, a nucleotide sequence encoding an amino acid sequence recognizable by a protein kinase will be designated a "PK nucleotide sequence". Similarly, the amino acid sequence recognizable by a protein kinase will be designated "PK amino acid sequence". A PK nucleotide sequence can be introduced into any selected nucleotide sequence, specifically a nucleotide sequence encoding the amino acids of a protein, polypeptide, or peptide. Preferably, the PK sequence is inserted into or joined to a nucleotide sequence which codes for a protein in such a manner so that the protein product is biologically, biochemically or otherwise active. The term "biologically active" is used throughout in this generic sense. Thus, the modified protein will be phosphorylatable and still be biologically active. Additionally, the phosphorylated modified protein should also be appropriately active for the use desired. In some cases, the phosphorylated protein could be inactive biologically, yet still be useful in a radioimmunoassay, for example.

It is only necessary in accordance with the invention that there be incorporated that much of the amino acid consensus sequence that will contain or be the site for the phosphorylation. Likewise at the DNA level, it is only necessary that the codons be provided which code for the amino acids which will be recognized by the kinase. Lack of absolute substrate specificity which has been reported to be a general property of protein kinases (21, 100), may provide further latitude in the selection of nucleotide sequences, or the codons coding for the recognized amino acids.

By "incorporates", "incorporation" or "insertion" or equivalent term, it is not necessary that the amino acid recognition sequence (or the nucleotide sequence coding for that amino acid sequence (or part thereof) be actually within the desired protein (or the nucleotide sequence coding therefor); it is sufficient that it be part of the modified protein, or part of the nucleotide sequence (or of part thereof) coding for the modified product.

A specific embodiment of the concept of the invention is a genetically engineered Hu-IFN- α A of the invention which, unlike the known Hu-IFN- α A (native, natural or genetically engineered), is phosphorylatable and the phosphorylated modified Hu-IFN- α . The phosphorylated Hu-IFN- α -P has retained biological activity with high radio-specific activity (2,000-12,000 Ci/mmol). Phosphorylation was performed with the catalytic bovine heart kinase which is a cyclic AMP (cAMP)-dependent kinase. The amino acid recognition (or consensus) sequences for the cAMP-dependent protein kinase have been identified as Arg-Arg-Ala-Ser-Val and Arg-Arg-Ala-Ser-Leu and Arg-Arg-Ala-Ser-Val-Ala amongst others (20, 21). In general cAMP-dependent protein kinases recognize the sequence Arg-Arg-Xxx-Ser-Xxx (21). The amino acid and nucleotide sequences of Hu-IFN- α A and their corresponding coding sequence have been reported (23-25).

Serine has been shown to be the target amino acid for phosphorylation for both the murine (5) and human (4, 10) IFN-gamma. It has also been shown that in Hu-IFN-gamma there are two serines capable of phosphorylation by the cAMP-dependent protein kinase, serines 132 and 142 (5, 10).

In accordance with the invention, in three specific illustrations of phosphorylated Hu-IFN- α (herein designated as Hu-IFN- α -P1, -P2, and -P3), the carboxyl-terminal amino acids were modified to contain the phosphorylation sites, and the nucleotide sequences encoding the putative phosphorylation sites were constructed.

The modified interferons were constructed with the use of oligodeoxyribonucleotides designed to produce insertions and substitutions at the carboxyl terminus of Hu-IFN- α A by site-specific mutagenesis procedures (described in the literature and also further described below) with the appropriate DNA sequences inserted into phage M13mp19. The general methods used for synthesizing a Hu-IFN- α with a modified carboxyl terminus in *E. coli* are further described in the following references (15, 26, 27).

It is not necessary that the amino acids at the carboxyl terminus be the ones that are modified; the modification can

be at the amino terminal end of the sequence, and correspondingly can be towards the 5' end of the nucleotide sequence; the modification can be virtually at any position of the protein sequence (and corresponding coding region) as long as the phosphorylation site is recognizable by the kinase in the intact protein.

As illustrated above with respect to Hu-IFN- α , the introduction into nucleotide sequences of one or more putative sites encoding amino acid sequences recognizable by an enzyme like a protein kinase opens wide and important new possibilities for modifying proteins in which such sites are absent, or inaccessible, or into which it is desired to introduce such additional sites.

Amongst the protein kinases there are known cyclic nucleotide-dependent protein kinases which catalyze the transfer of the gamma-phosphate of ATP to serine and/or threonine hydroxyl groups of acceptor protein substrates, calcium-dependent protein kinases, tyrosine-specific protein kinases, cyclic nucleotide- and calcium-independent protein kinases. The invention contemplates the use of any of the protein kinases, the nucleotide sequences which code for one or more of the corresponding putative sites for phosphorylation and the modified protein containing the amino acid sequences recognizable by this selected protein kinase, and the phosphorylated protein containing the radio-label. Such label can be phosphorus, or sulfur, or other groups discussed herein. Also, it need not be a radio-label. It is sufficient that it be an "identifiable" label.

Broadly considered, the invention contemplates the phosphorylation of various proteins by protein kinases after the introduction of phosphorylatable sites. Thus, the wide applicability of the invention in providing appropriate sites for phosphorylation in proteins otherwise not phosphorylatable can be recognized.

Throughout this description of the invention, the terms "phosphorylatable" (or "phosphorylated") are used. Reference is also being made to the analogs of the phosphate donor ATP, such as the gamma thiophosphate analog. Through this description the term "phosphorylation" or like term is intended to be generic to include "thiophosphorylation" where for instance, ^{31}P , ^{32}P , or ^{33}P is replaced by ^{35}S or sulfur analogs.

As discussed herein, kinases have been reported to catalyze phosphorylation not only at serine but also at threonine or tyrosine (21, 22, 64). Thus, it is within the scope of the invention to construct the nucleotide sequence which codes for the putative site(s) of recognition for the kinase selected and, in a manner analogous to that described herein, construct the replicable expression vehicle, introduce it into an appropriate host which in turn will express the modified phosphorylatable protein. In appropriate cases, the modified phosphorylatable protein is exogenous and mature. The phosphorylatable proteins can then be phosphorylated. Of course likewise, analogs of phosphorylated proteins -- e.g., sulfur-labelled -- can be made.

The nucleotide sequence encoding the putative phosphorylation site in the case of modified Hu-IFN- α which is recognized by the cAMP-dependent protein kinase is created by the oligodeoxyribonucleotide-directed insertion on the level of DNA as shown in FIG. 9.

Construction of the Modified Interferons.

The construction of expression plasmids for Hu-IFN- α containing phosphorylation sites was carried out as follows. The EcoRI-PstI fragment from pFLrA that contained the coding sequence for Hu-IFN- α A (23, 24) was inserted into the EcoRI-PstI site of M13mp19 to form M13BL27 that was used as the template for site-specific alterations as shown in FIG. 9. To construct the coding sequences for Hu-IFN- α A-P1 and P2, two oligodeoxyribonucleotides were synthesized to anneal to M13BL27 with the formation of a loop that permits the insertion of nucleotides to generate a coding sequence for a phosphorylation site at the COOH-terminal end (boxed residues, FIG. 8). The oligonucleotides used to prepare Hu-IFN- α A-P1 and -P2, respectively, were: AGT-TTA-AGA-AGT-AAG-AGA-AGG-GCA-AGT-GTT-GCA-TGA-AAA-CTG-CTT-CAA; and ACA-AAC-TTG-CAA-AGA-AGT-TTA-AGA-AGG-GCA-AGT-TTA-GCA-TGA-AAA-CTG-CTT-CAA. The underlined nucleotides are homologous with the coding sequence and 3' non-coding nucleotide sequence of the cDNA encoding Hu-IFN- α A; the nucleotides not underlined produce a loop for the insertion of additional residues for P1 and P2. A site-specific mutation as well as an insertion was introduced with the P2 oligonucleotide. After annealing of the oligonucleotide to single-stranded DNA from M13BL27, the second strand was synthesized and then cut with restriction endonucleases EcoRI-PstI. The EcoRI-PstI fragment obtained was then reinserted into the EcoRI-PstI site of phage M13mp19 as shown in Figure 9, and then E. coli were transformed with the duplex DNA. Incomplete duplexes (upper right of FIG. 9) do not yield the PstI-EcoRI duplex fragment. This excision and religation step was introduced to increase the efficiency of the site-specific mutations providing an overall yield of about 40% positive clones. RF DNA samples, prepared from individual phage M13 plaques, were screened for the presence of the inserted EcoRI-PstI fragment. Positive clones (i.e., those with insertions; about 75-90% of the plaques) were sequenced by the known Sanger dideoxynucleotide procedure (68) to identify the proper mutated recombinant and to confirm the sequence. By this procedure, about 50% of the transformants sequenced contained the mutated coding sequence with the phosphorylation site. The EcoRI-PstI fragments were then excised from the respective RF DNA preparations from the phages (M13BL28, M13BL29 and M13BL30) and religated into the EcoRI-PstI site of pRC23t to yield the expression vectors pBL281, pBL291 and pBL301 as shown in Figure 9. The two EcoRI-PstI fragments originating from pRC23t were obtained by

restriction endonuclease digestion of an expression plasmid for Mu-IFN- β that contained the trp terminator just downstream from the IFN- β coding sequence.

During the construction of the Hu-IFN- α A-P1 expression vector, a clone was isolated with a duplication of the -P1 oligonucleotide, a single nucleotide deletion of one A of codon 164 for Lys, and a second deletion of 11 3'-terminal residues (AA-CTG-GTT-CAA) from the downstream -P1 oligonucleotide. This series of steps generated an in-phase coding sequence for a new phosphorylation site on a slightly larger molecule designated Hu-IFN- α A-P3 (FIGS. 8 and 9).

The general recombinant DNA procedures employed herein have been described elsewhere (15, 26, 27). These methods are incorporated herein by reference. The use of vectors containing the phage lambda P_L promoter for cloning of recombinants and expression of proteins has been described by a number of laboratories (12, 48, 49, 70-77). A variety of E. coli strains lysogenic for wild type and mutant phage lambda or containing plasmids encoding the phage lambda repressor (12, 15, 48, 49, 70-78) have been used for growth of plasmids and/or expression of plasmid encoded proteins. For example, the following E. coli strains among others have been used for replication of plasmids and/or for expression of proteins under control of the phage lambda P_L promoter: E. coli 294 (75; ATCC #31977), AR 13 (77), AR58 (77), AR68 (75), AR120 (77, 74), C600 (49, 71; ATCC #33766), N99 (72; ATCC #33956), RR1 (66, 70; ATCC #31343), W3110 (77). With E. coli strains carrying a temperature-sensitive repressor (cl857), the cells can be grown at 30°C to prepare plasmids; upon shifting the temperature to 42°C, the repressor is inactivated and the DNA sequence under control of the phage lambda P_L promoter expressed (49, 66, 70-73, 75). Thus, cells can be grown to a high density at 30°C, then induced to express the protein of interest at 42°C. Alternatively, with the use of wild type phage lambda repressor (cl-), nalidixic acid can be used to induce expression of genes under control of the phage lambda P_L promoter (77). In the work involving this invention, it has been found desirable to use the temperature sensitive repressor (clts857) to regulate expression of the genes under control of the phage lambda P_L promoter. Although it was not found necessary to use protease deficient strains to achieve high production of recombinant proteins in E. coli (49, 70, 74, 77), strains lacking proteases (79, 80, 81) might prove beneficial in some cases (75, 76).

Construction and identification of bacterial plasmids containing the nucleotide sequence for Hu-IFN- α A is described in references 23 and 24. The reports describe the isolation of recombinant plasmids carrying DNA sequences corresponding to Hu-IFN- α species. The E. coli strain K-12 derivative RR1 can be considered a useful host in the present invention.

Site-specific mutagenesis which is well-suited for the present purposes is described in the following citations (15, 26, 27 and 43 at unit 8, supplement 2), which are incorporated herein by reference. The background of the method is reviewed in reference 44. A description in the patent literature is found in U.S. Patent No. 4,751,077, which is also incorporated herein by reference.

In accordance with the invention, the nucleotide sequence coding for the amino acid consensus sequence (which will contain the putative phosphorylation site) may be inserted anywhere into the nucleotide sequence of Hu-IFN- α A. This can be observed from the oligonucleotide sequences shown in FIG. 8. However, the insertion of the nucleotide sequence is preferably made at a site in the nucleotide sequence encoding Hu-IFN- α A so as to minimize an undesirable effect on the biological activity of the resultant recombinant protein, when such biological activity is critical.

The Expression of Hu-IFN- α A-P by Bacteria

The expression of the modified interferons in E. coli transformed with the expression plasmids encoding the Hu-IFN- α A-P proteins and the purification of Hu-IFN- α A-P proteins were carried out as follows. Each of the three above-named vectors was introduced into E. coli cells containing the plasmid pRK248clts857. E. coli AR68 cells and E. coli RR1 cells containing the compatible plasmid pRK248clts857 harboring each of the pBL281, pBL291 and pBL301 plasmids expressed Hu-IFN- α A-P1, -P2, and -P3, respectively, under control of the P_L promoter and trp terminator at 42°C. The yields of Hu-IFN- α A-P1, -P2, and -P3 in E. coli AR68(pRK248) were higher than that in E. coli RR1(pRK248) harboring the same plasmids. The products expressed in E. coli AR68 at 42°C were used to purify the Hu-IFN- α A-P1, -P2, and -P3 proteins by immunoaffinity chromatography as described further below. The data in Table 1 (below) show that greater than 50% of the antiviral activity was recovered. Hu-IFN- α A-P1, -P2, and -P3 were purified to a specific activity of 1.2×10^8 , 1.1×10^8 and 1.5×10^8 units per mg protein, respectively.

Plasmid pBL281 is deposited with the ATCC under Accession number 40509, and designated as pHu-IFN- α A-P1.

Table 1

Purification of Hu-IFN- α -P Proteins						
Human IFN- α A	Step	Antiviral activity		Protein	Specific Activity units/mg	Purification factor
		Total units	Yield (%)	mg		
α A-P1	1	1.6×10^7	100	32.6	4.9×10^5	1
	2	1.8×10^7	113	18.4	9.9×10^5	2
	3	1.1×10^7	71	0.09	1.2×10^8	246
α A-P2	1	1.6×10^7	100	34.3	4.7×10^5	1
	2	1.2×10^7	75	12.9	9.3×10^5	2
	3	0.9×10^7	56	0.08	1.1×10^8	232
α A-P3	1	1.6×10^7	100	33.0	4.8×10^5	1
	2	1.8×10^7	113	15.0	1.2×10^6	2.5
	3	2.6×10^7	163	0.18	1.5×10^8	312

The characterization of Hu-IFN- α A-P and its ^{32}P -labelled product was carried out as follows. The purified Hu-IFN- α A-P1, -P2, and -P3 proteins were analyzed by SDS-polyacrylamide gel electrophoresis in the presence of 2-mercaptoethanol (FIG. 10). A single band was observed on the silver stained gel with Hu-IFN- α A-P1 or -P2. Hu-IFN- α A-P2 migrated slightly faster than Hu-IFN- α A-P1 and a little slower than Hu-IFN- α A. Hu-IFN- α A-P3 yielded two bands, the slower migrating band being the intact molecule.

Hu-IFN- α A-P1, -P2, and -P3 were phosphorylated by the cAMP-dependent protein kinase with [γ - ^{32}P]ATP to a specific radioactivity of 2,000-12,000 Ci/mmol. Following phosphorylation, [^{32}P]Hu-IFN- α A-P1 and -P2 migrate on SDS-polyacrylamide gel electrophoresis in the presence of 2-mercaptoethanol as a single band with an apparent molecular weight of 19,000-20,000 (FIG. 11) corresponding to the same positions as the silver stained unlabelled band. The labelled Hu-IFN- α A-P3 migrates slower than the -P1 or -P2 products as expected. Since the faster migrating P3 band is unlabelled (cf. FIGS. 10 and 11), it is likely that the COOH-terminal extension of -P3 that contains the phosphorylation site is trimmed from the full length product to yield the faster migrating form. Unlabelled Hu-IFN- α A and [^{125}I]Hu-IFN- α A were used as molecular weight standards and controls (FIGS. 10 and 11).

Antiviral Activity of the New Interferons.

The effect of phosphorylation on the antiviral activity of Hu-IFN- α A-P1, -P2, and -P3 was determined in a parallel experiment. Phosphorylation, it was found, has little or no effect on the antiviral activity of the Hu-IFN- α A-P1, -P2, and -P3 measured with both bovine MDBK cells and human WISH cells.

Table 2, below shows the antiviral activity of non-phosphorylated and phosphorylated interferon alpha proteins.

Table 2

Effect of Phosphorylation on the Antiviral Activity of Phosphorylated Human Interferon Alpha Proteins			
Human IFN- α A	[γ - ^{32}P]ATP	Antiviral Activity	
		MDBK	WISH
α A-P1	-	1.6×10^6	4.0×10^6
	+	1.6×10^6	2.4×10^6
α A-P2	-	1.6×10^6	8.8×10^6
	+	0.9×10^6	8.8×10^6
α A-P3	-	6.4×10^6	4.8×10^6
	+	4.8×10^6	3.2×10^6

Ability to Bind to Receptors.

The ability of [^{32}P]Hu-IFN- α A-P to bind to receptors was shown as follows. [^{32}P]Hu-IFN- α A-P1 and -P2 bind to

bovine MDBK and human Daudi cells (FIG. 12) with the specific binding approaching saturation at higher concentrations. Scatchard analysis of the data yielded the following estimates. The bovine MDBK cells contain approximately 3,800 and 9,450 receptors per cell calculated from the binding of [32 P]Hu-IFN- α -P1 and -P2, respectively; Daudi cells, approximately 1,650 and 4,900 receptors per cell. Dissociation constants (K_d) were calculated to be 1.4×10^{-10} M for the binding of [32 P]Hu-IFN- α -P1 to both the human and bovine cells; and 3.5×10^{-10} M and 2.2×10^{-10} M for the binding of [32 P]Hu-IFN- α -P2 to human and bovine cells, respectively. Similar results are obtained with [32 P]Hu-IFN- α -P3.

All these phosphorylated [32 P]Hu-IFN- α -P derivatives bind to the Hu-IFN- α/β receptor because their binding to Daudi cells was competitively blocked by Hu-IFN- α and Hu-IFN- β , but not by Hu-IFN- γ .

Crosslinking of [32 P]Hu-IFN- α -P to the Receptors.

The crosslinking of [32 P]Hu-IFN- α -P to the receptors was carried out as follows. [32 P]Hu-IFN- α -P1, -P2, and -P3 were each covalently crosslinked to the receptors by reaction with disuccinimidyl suberate after binding to the cells (FIG. 13). The radioactive complexes migrate as several bands with molecular weights of 100K-200K from the Daudi cells or as a broad band with an apparent molecular weight of about 150K from the MDBK cells upon analysis by SDS-polyacrylamide gel electrophoresis. The crosslinked complexes of the receptors on the cells with [32 P]Hu-IFN- α -P1, -P2, and -P3 appear to be the same on SDS-polyacrylamide gels, but differ from the crosslinked complexes of [32 P]Hu-IFN- γ formed with human Daudi cells. Neither the crosslinked complexes nor the free [32 P]Hu-IFN- α -P1, -P2, and -P3 are seen if excess non-radioactive Hu-IFN- α is included during the binding reaction (FIGS. 12 and 13).

Hu-IFN- β Containing A Phosphorylation Site.

Likewise nucleotide sequences coding for a Hu-IFN- β -like protein, which sequence contains a putative phosphorylation site recognizable by the cAMP-dependent protein kinase, is prepared.

The following procedure generates a modified Hu-IFN- β containing a site recognizable by the cAMP-dependent protein kinase from bovine heart. The PstI-BglII 363 bp fragment from the cDNA encoding Hu-IFN- β is excised from the expression vector pFIFrp69 (104). The 363 bp fragment is inserted into the PstI and XmaI sites of phage M13mp18. First the PstI end of the PstI-BglII fragment is ligated to the dsDNA of M13mp18 cut with restriction endonucleases PstI and XmaI. The BglII end of the fragment is then ligated to the M13mp18 vector with the use of a BglII-XmaI linker:

GATCTGCGCGCGC

ACGCGCGCGGGCC

which reconstructs BglII and XmaI sites. Since there is no BglII site in the polylinker region of M13mp18, this BglII-XmaI linker is used to preserve the BglII site and permit cutting with XmaI for preparation of one of the intermediate recombinants (see below and analogous construction in Fig. 8). The M13mp18 containing the 363 bp fragment from the 5' end of the coding region of Hu-IFN- β formed M13-A that is used as a template for site specific mutation as follows. The site specific insertion is made similarly to that described above for construction of Hu-IFN- α -P1. To construct the coding sequence for Hu-IFN- β -P, the modified Hu-IFN- β , a oligodeoxyribonucleotide(CTT-ACA-GGT-TAC-CTC-CGA-AGG-GCA-AGT-GTT-GCA-TGA-AGA-TCT-GCG-CGC-GCC-CGG) is synthesized to anneal to M13-A with the formation of a loop that would permit the insertion of nucleotides to generate a coding sequence for a phosphorylation site at the COOH-terminal end. The underlined residues of the oligonucleotide shown above are homologous with the nucleotides of the phage M13-A that contains the cDNA fragment encoding Hu-IFN- β . A comparison of Hu-IFN- β and Hu-IFN- β -P is shown below:

160

165

Hu-IFN- β Leu-Thr-Gly-Tyr-Leu-Arg-Asn-END

CTT-ACA-GGT-TAC-CTC-CGA-AAC-TGA

160

165

170

Hu-IFN- β -P Ieu-Thr-Gly-Tyr-Leu-Arg-Arg-Ala-Ser-Val-Ala-END

CTT-ACA-GGT-TAC-CTC-CGA-AGG-GCA-AGT-GTT-GCA-TGA

The underlined residues of the modified Hu-IFN- β -P from 165-170 represent the cAMP-dependent protein kinase recognition site. The nucleotides of the oligonucleotide above that are not underlined produce a loop for the insertion of residues 166-170 of Hu-IFN- β -P. After annealing of the oligonucleotide to single-stranded DNA from M13-A, the second strand is synthesized and then cut with restriction endonucleases PstI and XmaI. The resultant PstI-XmaI fragment is then reinserted into the PstI-XmaI site of phage M13mp18 as shown analogously in Fig. 8 and then E. coli are transformed with the duplex DNA. This excision and religation step is introduced to increase the efficiency of the site-specific mutations. RF DNA preparations from individual phage M13 plaques are screened for the presence of the inserted PstI-XmaI fragment. Positive clones (i.e. those with insertions) are sequenced by the Sanger dideoxynucleotide procedure (68) to identify the proper mutated recombinant and to confirm the sequence. By this procedure, several transformants sequenced contain the mutated coding sequence with the phosphorylation site (M13- β). The PstI-BglII fragment is then excised from the RF DNA from the phage M13-B and religated into the PstI-BglII site of pFIFtrp69 to yield the expression vector similar to the original vector. The general recombinant DNA procedures employed have been described elsewhere (15, 26, 27, 43).

The phosphorylatable modified interferon is expressible as described (104), and can be purified following the procedure for the corresponding Hu-IFN- β (65).

The following illustrative examples are not intended to limit the invention in any manner whatsoever.

ILLUSTRATIVE EXAMPLES

Bacterial Strains, Enzymes and Chemicals. E. coli RR1 containing the plasmid pRK248cls857 with the temperature-sensitive repressor of the phage lambda P_L promoter, was obtained from Robert Crowl (12). The plasmid pPRK248cls857 was introduced into E. coli AR68 (75) by transformation.

Restriction endonucleases and polynucleotide kinase were from New England BioLabs. The buffer conditions used were described by the manufacturer. The Klenow fragment of Escherichia coli DNA polymerase I and T4 DNA ligase were from International Biotechnologies, Inc.; calf intestinal phosphatase (c.i.p.) was from Boehringer-Mannheim Biochemicals. Ligation reactions were carried out in the presence of low melting point agarose gel (15, 66).

The catalytic subunit of cAMP-dependent protein kinase from the bovine heart muscle with a specific activity of 20,000 units/ml was obtained from Sigma. [γ - 32 P]ATP with specific radioactivity of 5,000 Ci/mmol was obtained from Amersham; dithiothreitol (DTT), from Bethesda Research Laboratories; bovine serum albumin (BSA), from Miles Laboratories; acrylamide and N,N'-methylenebisacrylamide, from International Biotechnologies, Inc. (IBI); sodium dodecylsulfate (SDS), from Sigma; and disuccinimidyl suberate (DSS), from Pierce Chemical Co.

Interferon and Protein Assays. Interferon activity was determined by a cytopathic effect inhibition assay with vesicular stomatitis virus and bovine MDBK cells (13). All interferon titers are expressed in reference units/ml calibrated against the reference standard for human leukocyte interferon (G-023-901-527) obtained from the Antiviral Substances Program of the National Institute of Allergy and Infectious Diseases, National Institutes of Health, Bethesda, MD. Protein concentration was determined by the procedure of Bradford (14) with bovine serum albumin as a standard.

The following illustrates the preparation of the fused-hybrid Hu-IFN- α /gamma.

Synthesis and Phosphorylation of Oligonucleotide.

The oligodeoxyribonucleotide CTGACTCCTTTTCGCTTTTCCTTACTTCTTAAC which was used for oligonucleotide-directed mutagenesis and hybridization screening was synthesized and phosphorylated as described (66). After the phosphorylation reaction, the reaction mixture was diluted with 3.5 ml of 6 x SSC (0.9 M NaCl, 90 mM sodium citrate)

and used directly for screening by hybridization.

Oligonucleotide-directed Deletion. About 100 pmol of phosphorylated oligomer were annealed with 1 pmol of ssDNA template in 10 μ l of 30 mM Tris-HCl, pH 7.5, 10 mM MgCl₂ at 80°C for 5 min. and gradually cooled to 30°C for 30 min., then put on ice. The volume was adjusted to 20 μ l of 30 mM Tris-HCl, pH 7.5, 10 mM MgCl₂, 0.5 mM dATP, 0.5 mM dCTP, 0.5 mM dGTP, 0.5 mM dTTP, 1 mM ATP, 10 mM dithiothreitol (DTT). The oligomer was extended with 2.5 units of the Klenow fragment of DNA polymerase I in the presence of 6 units of T4 DNA ligase at 16°C for 24 hr. The reaction mixture was then extracted once with an equal volume of phenol and chloroform (1:1) and precipitated with 1/10 volume of 2.5 M sodium acetate and 2 volumes of 95% ethanol. After washing with 75% ethanol, DNAs were digested with *Spe*I for the enrichment of the mutated DNA before transformation of RR1(λ clt₈₅₇). For screening, the colony hybridization method was used as described (58, 67). DNA sequencing was performed by the dideoxy sequencing method (68) after the 600 bp *Hinc*II/*Pst*I fragment from M13mp18/Hu-IFN- α /gamma was inserted into the *Hinc*II and *Pst*I site of M13mp19.

Preparation and Purification of Hu-IFN- α /gamma. *E. coli* RR1(pRK248clt₈₅₇) cells harboring Hu-IFN- α /gamma were grown in LB broth at 30°C overnight. An overnight culture of 40 ml was diluted to 2,000 ml of M9 medium containing 0.4% of casamino acids, 50 μ g/ml ampicillin (Ap) and 25 μ g/ml tetracycline. Bacteria were grown at 30°C until the cell density at 600 nm reached 1.0, at which time the cells were transferred to 42°C for 2-3 hrs. The cells were collected by centrifugation at 7,000 rpm for 10 min. and stored at -20°C until used.

For purification, all steps were carried out at 4°C. Frozen cells (10 g) were suspended in 3 volumes (40 ml) of 7 M guanidine hydrochloride in 25 mM Tris-HCl, pH 7.5. The suspension was mixed for 1 hr. and centrifuged at 17,000 rpm for 30 min. The supernatant was diluted into 10 volumes of phosphate buffered saline (PBS). Solid ammonium sulfate was added to 65% saturation with vigorous stirring. The suspension was kept at 4°C for 2 hrs. and then centrifuged at 10,000 rpm for 20 min. The pellet was suspended in 20 ml of PBS and dialyzed extensively against PBS. The suspension obtained was centrifuged at 10,000 rpm for 20 min., and the supernatant was mixed with 0.3 ml of monoclonal antibody to Hu-IFN- α (LI-8) coupled to Affi-gel 10 (16, 41) for 1 hr. The mixture was then loaded into a 1-ml disposable syringe. After washing with four column volumes (0.3 ml) of 0.5 M NaCl, 25 mM Tris-HCl, pH 7.5, the column was rinsed with four column volumes of 0.15 M NaCl and then eluted with four column volumes of 0.2 N acetic acid and 0.15 M NaCl, pH 2.5. Antiviral activity was eluted in the first two fractions. It was measured by a cytopathic effect inhibition assay on bovine MDBK cells (13). The concentration of protein was determined by the procedure of Lowry (69) or Bradford (14) with bovine serum albumin as a standard. The concentrations obtained by the two procedures were in agreement. The procedure yielded 240 μ g of Hu-IFN- α /gamma with a specific activity of 4×10^6 units/mg.

Preparation of Hu-IFN- α /gamma protein from *E. coli* AR68(pRK248clt₈₅₇) harboring the expression plasmid pXZ-8 and its purification by immunoaffinity chromatography with monoclonal antibody against Hu-IFN- α is as described below for the Hu-IFN- α -P proteins. The yield from 50 ml of bacterial culture was 388 μ g of Hu-IFN- α /gamma with a specific activity of 2.3×10^7 units/mg. The procedure of purification by immunoaffinity chromatography with monoclonal antibody against the COOH-terminal sequence of Hu-IFN-gamma is the same as above. Elution with acid yielded 281 μ g of Hu-IFN- α /gamma with a specific activity of 1.3×10^7 units/mg from 50 ml of bacterial culture.

Phosphorylation of Hu-IFN- α /gamma. Hu-IFN- α /gamma was phosphorylated as described for Hu-IFN-gamma (3, 6) with some minor modifications. About 1 μ g of Hu-IFN- α /gamma was incubated at 30°C for 60 min. with 0.5 mCi of [γ -³²P]ATP (>5,000 Ci/mmol, Amersham Corp.) and 10 units of the catalytic subunit of bovine heart cAMP-dependent protein kinase (Sigma) in 30 μ l of 20 mM Tris-HCl, pH 7.4, 100 mM NaCl, 12 mM MgCl₂, and 3 mM DTT, then cooled on ice to stop the reaction. After addition of 0.3 ml of 5 mg/ml bovine serum albumin in 10 mM sodium pyrophosphate, pH 6.7, the reaction mixture was dialyzed extensively against 10 mM sodium pyrophosphate, pH 6.7, at 4°C. Incorporation of radioactivity into Hu-IFN- α /gamma was measured with a liquid scintillation spectrometer after precipitation of the protein with trichloroacetic acid (82).

Binding of Hu-IFN- α /gamma to cells. Bovine MDBK cells were used for binding studies. MDBK cells were grown to confluence in 6-well tissue culture plates in medium (Gibco F-11) supplemented with 10% fetal calf serum and 50 μ g/ml gentamicin. Medium was removed, and 1 ml of fresh medium was added into each well. After 20 min., [³²P]Hu-IFN- α /gamma was added in the absence or presence of 0.6 μ g unlabelled Hu-IFN- α (10^8 units/mg). Following incubation at room temperature for 1 hr., the plates were put on ice and each well was washed with three 1 ml volumes of cold PBS. Then 1 ml of 1% SDS was added to remove cells from the wells. Radioactivity was determined with a liquid scintillation counter by placing the 1 ml samples in 10 ml of a scintillation fluor (83). Binding of [³²P]Hu-IFN- α /gamma to the human Daudi cells is described below with the Hu-IFN- α -P proteins.

The following illustrates the construction of modified interferons by "insertion".

Construction of Modified Interferons. Because of certain limitations of the hybrid-fusion procedure and product as discussed above, an alternative construction was explored. It was then discovered that a putative phosphorylation site could be introduced into a nucleotide sequence of Hu-IFN- α .

The amino acid recognition or consensus sequences for the cAMP-dependent protein kinase have been identified as Arg-Arg-Ala-Ser-Val and Arg-Arg-Ala-Ser-Leu among others (20, 21). The amino acid sequence of Hu-IFN- α and

its corresponding coding sequence have been reported (23-25). Hu-IFN- α A as well as other interferons have been expressed in *E. coli* expression vectors under control of the *trp* and the phage lambda P_L promoter (see references 1, 24 and 25 for reviews and additional citations). The terminal nucleotides corresponding to the last ten carboxyl-terminal amino acids of Hu-IFN- α A are shown in FIG. 8 as well as sequences corresponding to modified molecules, Hu-IFN- α A-P1, -P2, and -P3, which contain putative phosphorylation sites. To construct these molecules oligodeoxyribonucleotides were synthesized to introduce the insertions and substitutions shown at the carboxyl terminus of Hu-IFN- α A (FIG. 8) by site-specific mutagenesis procedures with the appropriate DNA sequences inserted into phage M13mp19 (15, 26, 27). The phosphorylation sites in Hu-IFN- α A-P1, -P2, and -P3 (FIG. 8) recognized by the cAMP-dependent protein kinase were created by the oligodeoxyribonucleotide-directed insertion on the level of DNA as shown in FIG. 8.

The expression plasmids pBL281, pBL291 and pBL301, coding for Hu-IFN- α A-P1, -P2, and -P3, respectively, were constructed as outlined in FIG. 8. The sequences coding for the modified interferons were inserted into an expression vector under control of the phage lambda P_L promoter also as illustrated in FIG. 8. *E. coli* AR68 containing the compatible plasmid pRK248cIts857 was transformed with each of the expression plasmids encoding the Hu-IFN- α A-P proteins containing the COOH-terminal sequences shown in FIG. 8. The phosphorylatable modified interferons were expressed and purified as described further below.

Expression and Preparation of Hu-IFN- α A-P. *E. coli* RR1 (pRK248cIts857) cells harboring pBL281, pBL291 or pBL301 plasmids containing the Hu-IFN- α A-P1, -P2, and -P3 coding sequences were grown at 30°C overnight in M9CA medium (15) with the concentration some components modified as follows: 1% casamino acids, 1% glucose, 10 mM MgSO₄ and 1 mM CaCl₂, and containing 2 µg/ml thiamine, 50 µg/ml ampicillin and 12.5 µg/ml tetracycline. For expression of Hu-IFN- α A and modified proteins, 100 ml of M9CA medium was inoculated with 3-5 ml of an overnight culture. The bacteria were grown at 30°C until the cell density reached an optical density at 600 nm of 0.3-0.5 in 2-3 hours, at which time the culture was transferred to 42°C for an additional two hours. The bacterial cells were collected by centrifugation and lysed in 8 M guanidine hydrochloride and 50 mM Tris-HCl, pH 7.6, at 0°C for 10 min. The supernatant obtained after centrifugation at 14,000 rpm (SA-600 Sorvall rotor) for 30 min. was used to assay the antiviral activity or to purify the Hu-IFN- α A and modified interferons.

E. coli AR68(pRK248cIts857) cells harboring plasmids pBL281, pBL291, or pBL301 containing Hu-IFN- α A-P coding sequences were grown in LB medium (15) containing 50 µg/ml ampicillin and 12.5 µg/ml tetracycline at 32°C overnight. The overnight culture was diluted five-fold with fresh LB medium containing the same concentration of antibiotics as above and then grown at 32°C for 2-3 hours. For expression of the Hu-IFN- α A-P proteins, harvesting of cells, and preparation of supernatants the procedures described above for use with *E. coli* RR1 cells were used.

The purification of Hu-IFN- α A-P proteins were carried out as follows. All steps for purification of Hu-IFN- α A-P species were carried out at 4°C-8°C. Ten ml of the guanidine hydrochloride supernatant from 100 ml of the expressed culture were diluted tenfold with cold phosphate-buffered saline (PBS) and precipitated at 65% saturation of ammonium sulfate at 4°C overnight. The precipitate was collected by centrifugation at 10,000 rpm (Sorvall GSA rotor) for 20 min. at 5°C. The supernatant was decanted and saved. The residual pellet was dissolved again in 10 ml of cold PBS and the solution centrifuged as above. The combined supernatants of 30 ml were mixed with 1 ml of Affi-gel 10 to which monoclonal antibody LI-8 (against Hu-IFN- α) was linked (16, 41) and the suspension rocked at 4°C for 1 hour. The immunoabsorbent was loaded into the barrel of a 2 ml disposable syringe and washed with 20 column volumes or more of each of the following cold solutions sequentially (16): PBS, Buffer F (0.5 M NaCl, 25 mM Tris-HCl, pH 7.5, and 0.2% Triton X-100); and 0.15 M NaCl. Then the interferon was eluted with Buffer H (0.2 M acetic acid, 0.15 M NaCl, pH 2.6) and 0.4 ml fractions collected. The eluted fractions were neutralized with 1 M Tris base to pH 7.0 and the fractions of peak antiviral activity pooled.

The phosphorylation of Hu-IFN-gamma and Hu-IFN- α A-P proteins was carried out as follows: Hu-IFN- α A-P or Hu-IFN-gamma were labelled with [gamma-³²P]ATP and the cAMP-dependent protein kinase as described for Hu-IFN-gamma with some minor modifications (3, 6). About 0.25 to 0.65 µg of Hu-IFN-gamma or Hu-IFN- α A-P was incubated at 30°C for 1 hour with 0.25 mCi of [gamma-³²P]ATP (5,000 Ci/mmol, Amersham) and 7.5 units of the catalytic subunit of cAMP-dependent protein kinase in 30 µl containing components as previously reported (3, 6). The reaction mixture was then cooled in an ice bath, and, after the addition of 270 µl of 5 mg/ml bovine serum albumin in 10 mM sodium pyrophosphate (NaPPi), pH 6.7, was dialyzed extensively against 10 mM NaPPi at 4°C. The radioactivity associated with [³²P]Hu-IFN- α A-P was determined in a Beckman Model-LS3801 scintillation spectrometer. The phosphorylated Hu-IFN- α A-P was stored in liquid nitrogen in small volumes.

It has been reported (6) that the Hu-IFN-gamma phosphorylated with ³²P has a 100-fold higher specific radioactivity than reported for [¹²⁵I]IFN-gamma.

The phosphorylated interferons in accordance with the invention provides molecules with higher radio-specific activity than previously obtainable (1,000-12,000 Ci/mmol) with retention of biological activity. Thus, the phosphorylation site inserted into Hu-IFN- α A at the COOH terminus does not detrimentally affect the biological activity (antiviral activity) and can be effectively recognized by the cAMP-dependent protein kinase.

Further Hu-IFN- α A-P1 and -P2 are stable during purification and phosphorylation. The Hu-IFN- α A-P3, which as shown in FIG. 8 has an additional septidecylpeptide at the COOH terminus, degraded into at least two fragments.

It is evident that for some biological applications the phosphorylated modified interferons, and those labelled with phosphate analogs such as those containing S, should be stable in serum.

The binding of [32 P]Hu-IFN- α A-P1, -P2, and -P3 to bovine MDBK cells and human Daudi cells was performed. [32 P]Hu-IFN- α A-P1, -P2, and -P3 crosslinked to cells exhibited one complex of about 150K with the bovine MDBK cells and several complexes of 100-200K with the human Daudi cells.

The binding of [32 P]Hu-IFN- α A-P to cells was performed as follows: Confluent monolayers of bovine MDBK cells were trypsinized and 1 ml (1×10^6 cells) of the cell suspension in Dulbecco's modified Eagle's medium (Gibco) containing 10% inactivated fetal calf serum and 1% penicillin-streptomycin solution (Gibco) was added to each well of a 6-well plate. The cell monolayers reached confluence and approximately doubled on overnight incubation at 37°C at which time they were used to measure binding of [32 P]Hu-IFN- α A-P. For beginning the binding, the medium was removed, then 1 ml of fresh medium containing [32 P]Hu-IFN- α A-P at the indicated concentration was added into each well in the absence (-) or presence (+) of excess nonradioactive Hu-IFN- α A as a competitor (>500-fold more than [32 P]Hu-IFN- α A-P added). The plates were incubated with rocking at room temperature (24°C) for 60 min., after which they were placed on ice to cool. Each well was washed three times with 1 ml of cold PBS to remove the unbound radioactive ligand. After washing, 1.5 ml of 1% sodium dodecylsulfate in water was added to each well and, after dissolution of the cells and bound radioactivity, the entire 1.5 ml was counted in a Beckman Model LS3801 scintillation counter in 2 ml Hydrofluor scintillation fluid.

The binding of [32 P]Hu-IFN- α A-P to human Daudi cells was performed as described previously (17) with some modifications. Daudi cells were harvested by centrifugation at 1,000 x g for 10 min., washed twice with the growth medium (RPMI-1640, Gibco-H18, supplemented with 12.5 mM sodium HEPES, 10% fetal calf serum and 50 μ g/ml gentamicin) and resuspended in the medium to a concentration of 1×10^7 cells/ml. The binding of [32 P]Hu-IFN- α A-P at the indicated concentration to 1.25×10^6 Daudi cells in a total volume of 125 μ l was allowed to proceed in the absence or presence of non-radioactive Hu-IFN- α A as a competitor at room temperature (24°C) for 60 min., with gentle resuspension every 15 min. At the end of the 60 min. incubation period, 100 μ l of the cell suspension was layered onto a 300 μ l cushion of 10% sucrose in PBS in a sample cup and pelleted by centrifugation (Beckman Microfuge Type B) for 2 min. Tubes were frozen in a dry ice-ethanol bath, and then the tips of tubes containing the cell pellets were cut off and counted as above. The specific binding at a given concentration of [32 P]Hu-IFN- α A-P is defined as the difference in bound radioactivity between samples incubated in the absence (total) and presence (nonspecific) of excess non-radioactive Hu-IFN- α A.

The covalent crosslinking of [32 P]Hu-IFN- α A-P to the receptors was carried out as follows: A monolayer of bovine MDBK cells in 75-cm² tissue culture flasks was washed twice with Dulbecco's phosphate-buffered saline, trypsinized with 2 ml of trypsin-EDTA solution (1 x in phosphate-buffered saline, Gibco laboratories) at 37°C until the cells were released from the tissue culture flask. After addition of 10-20 ml of Dulbecco's modified Eagle's medium (Gibco Laboratories) containing 10% inactivated fetal calf serum and 1% penicillin-streptomycin (Gibco), cells were collected by centrifugation at 500 x g for 5-10 min. and resuspended in the same medium to a concentration of about 1×10^7 cells/ml. About 5×10^5 cpm of [32 P]Hu-IFN- α A-P (2,000-12,000 Ci/mmol) was added to 0.5 ml of cells with or without 1 μ g of non-radioactive Hu-IFN- α A as a competitor. After incubation with rocking at room temperature (24°C) for 1 hour, the cells were pelleted for 20 seconds at 14,000 rpm in an Eppendorf Microfuge, washed twice with 1 ml of cold PBS, then treated with a final concentration of 0.5 mM disuccinimidyl suberate (freshly prepared in dimethylsulfoxide) at 4°C for 20 min. as described (7). The crosslinking of [32 P]Hu-IFN- α A-P to cell receptor proteins was analyzed by sodium dodecylsulfate (SDS)-polyacrylamide gel electrophoresis (18) on 1.5-mm thick slab gels containing 8% acrylamide as described in detail previously (6, 7). The crosslinking thus was confirmed.

³⁵S-labelled proteins, like Hu-IFN- α A-P1, are obtainable in a manner similar to that described above.

SDS-polyacrylamide Gel Electrophoresis. The proteins, labelled products and covalent crosslinked complexes were analyzed by electrophoresis on the SDS-polyacrylamide slab gels of 1.0 or 1.5-mm thickness by the method of Laemmli (18). After electrophoresis, the proteins were stained with silver (19) or with Coomassie brilliant blue. Gels containing radioactive samples were dried under vacuum and autoradiographed at room temperature for the labelled ligands and at -170°C for the covalent crosslinked complexes with Kodak X-Omat film and intensifying screens.

Other Expression Vectors, Host Cells, Etc. While presently preferred procedures to express the modified interferons, to make various nucleotide sequences, and to transform specific hosts have been illustrated, it is evident that the invention is not in any way limited by these illustrations. Both eukaryotic and prokaryotic host cells may be used. Several procedures for the isolation of genes and expression of interferons in bacterial cells and heterologous cells are quite well-suited for production of modified interferons of the invention. For instance, several such methods describe in reference (1) Section VII the use of yeast vectors for production (Chapter 59) of and secretion (Chapter 60) of human interferons by yeast. Other microbial strains of *E. coli* may be used, or *Bacilli*, like *Bacillus subtilis*, *Salmonella typhimurium* (as disclosed in U.S. Patent No. 4,727,138) with plasmids that can replicate and express heterologous gene

sequences therein. Other expression vectors are illustrated in U.S. Patent No. 4,559,300, for instance. Numerous other promoter systems (than the one illustrated herein) can be used like the trp (104, 105) and lac (104, 106) promoters for example. All such references are included hereby by reference.

Likewise, the modified interferons can be produced from vertebrate cell cultures, for instance, a COS-7 line of monkey kidney fibroblasts can be used as the host for the production of the modified interferons with appropriate expression vectors (see Chapter 63 in reference 1, for example); other cell lines are suitable and are known. An example of the use of a retroviral based vector for expression in eukaryotic cells is given in Chapter 56 of reference 1. Many other examples of eukaryotic expression vectors have been described (see for example 50, 51, 84-90).

Vectors useful in the invention to replicate in a transformed host cell have a DNA segment containing a functional origin of replication (replicon). Plasmids and phage DNA by their very nature contain replicons facilitating replication in a host cell. The vector will have a DNA segment which conveys to a transformable host cell a property useful for selection of transformed cells from nontransformed cells. Any of a wide range of properties can be used for selection purposes. One of the most commonly used properties is antibiotic resistance, e.g., tetracycline resistance or ampicillin resistance.

The foregoing two elements generally are present in readily available and recognized cloning vectors. Examples of suitable cloning vectors are bacterial plasmids, such as plasmids from E. coli, including pBR322, pMB89, ColE1, pCR1; wider host range plasmids, including RP4; phage DNAs, such as lambda, and the like. Most, if not all, of the above-recognized vectors already carry the aforescribed two elements. Of course, as is known, in certain recombinants the DNA need not contain a replicon nor an attached marker.

Thus, any suitable expression vector may be used to express the modified interferons having putative phosphorylation sites in accordance with the invention.

In accordance with known procedures, the DNA comprises the regulating segments and the coding region. Thus, it is evident that the invention is not limited to the particular means of constructing genetic recombinants disclosed as illustrations and that one of average skill in the art would without undue experimentation adapt, change or select the procedures best suited to his objective.

For techniques and additional materials (vectors, host systems, plasmids, enzymes used in molecular cloning, synthesis and cloning of cDNA, introduction of plasmid and bacteriophage lambda DNA into E. coli, etc.), see (15) and (43), incorporated herein by reference.

ADDITIONAL GENERAL OBSERVATIONS

As has been described above, Hu-IFN- α A and Hu-IFN- β cannot be phosphorylated (by the cAMP-dependent bovine heart kinase) whereas it has been shown that Hu-IFN-gamma and the corresponding Ra- and Mu-IFN-gamma are amenable to phosphorylation without modification of the DNA-sequence (5, 11).

Thus, in accordance with the invention, additional phosphorylation sites can be introduced into Hu-IFN-gamma to provide interferon proteins that can be labelled to higher radio-specific activities than proteins, e.g., interferons, with only a single phosphorylation site.

Thus, it is within the contemplation of the invention to increase the number of sites (phosphorylation, thiophosphorylation) in proteins which already have one (or a larger number) of such sites.

The origin of the phosphorylatable nucleotide sequence can furthermore be varied. Instead of using the PK nucleotide sequence corresponding to Hu-IFN-gamma, there may be used a PK sequence corresponding to any other interferon (such as murine interferon), or for that matter the PK nucleotide sequence is derivable from any other nucleotide sequence encoding a protein known to be phosphorylatable. In this manner highly interesting phosphorylatable proteins (and phosphorylated proteins) can be made available for practical applications.

Furthermore, it is within the scope of the invention to radio-label proteins, like Hu-IFN- α A and - β which have not been phosphorylatable heretofore with a radioactive label which has advantages over iodine labelling. Accordingly, it is within the scope of the invention to use the sulfur analogs of the radioactive ATP wherein sulfur is substituted for the phosphorus. For instance, the gamma ^{35}S analog of ATP could be incorporated into the protein at the appropriate recognition site. Thus, the invention contemplates the use of the isotopes of sulfur and phosphorus, like ^{35}S , ^{38}S , ^{31}P , ^{32}P , and ^{33}P . Some of these isotopes have not yet been widely used because they are less readily available or because of their respective half-life. Lists of isotopes are of course available. Any isotope which can be introduced into a compound to be labelled is contemplated by this invention.

From Table 3 shown below, it can be observed that ^{38}S with a half-life of 2.87 hours and an energy of 1.1 Mev may well be an ideal source for irradiation of tumors and other tissues when radiation is deemed appropriate. Proteins phosphorylated with such an isotope with a short half-life of 2.9 hours would have a specific activity about one hundred times the specific activity that is obtainable with ^{32}P . The short half-life of the isotope also has the advantage that it is eliminated from the patient within a few days. Furthermore, the ability to generate specific activities one-hundred fold that of ^{32}P permits the use of smaller dosages of the proteins so that antigenic side-effects (that is, antibody production to these proteins) is minimized. The ^{33}P phosphate produces a β particle with an energy approximately one-seventh that of ^{32}P .

The isotope has a half-life of 24.2 days so that a protein phosphorylated with a phosphate analog containing ^{33}P would have approximately one-half the specific activity of the analog labelled with ^{32}P (with a 14.2 day half-life). The ^{33}P radiation would affect less of the area surrounding the tumor cells than the derivatives containing ^{32}P . Thus, the use of the products of the invention can bring about important beneficial advantages.

ADDITIONAL APPLICATIONS AND USES OF MODIFIED PROTEINS

The interferons modified in accordance with the invention by the presence of one or more phosphorylated groups -- or analogs thereof like sulfur -- and proteins so modified, have numerous applications and uses in the biological, medical, biomedical (including therapeutic and diagnostic), and other sciences.

It is contemplated that modified proteins in accordance with the invention can have additional specific uses. A few illustrations of such uses are described below.

1. Pharmacokinetics of proteins.

It is often useful to follow the fate of injected proteins in animals and patients. It is shown below that the phosphorus attached to some of these proteins is relatively stable in human and fetal calf serum; thus the pharmacokinetics of proteins can be conveniently studied. Thus, phosphorylated proteins are especially well-suited for such applications.

For uses of the phosphorylated proteins or analogs of the invention where the protein is expected to be in contact with human or animal serum, it is necessary that the protein derivative be stable in human or animal serum. The derivative protein should be stable in the serum of the species in which the pharmacokinetic studies (or application) are to be carried out, or in a serum equivalent, i.e., from the biological point of view, to the serum of the species on which the work is to be performed.

For instance, in the work described above, the phosphate linked to Hu-IFN- α A-P1, -P2, and -P3 is stable in fetal calf serum at 37°C. In the presence of human serum, the phosphate linked to Hu-IFN- α A-P1 and to Hu-IFN- α A-P3 was stable at 37°C, but the phosphate linked to Hu-IFN- α A-P2 was labile. After 6 and 16 hours at 37°C approximately 62% and 74% of the phosphate was hydrolyzed from [^{32}P]Hu-IFN- α A-P2. Thus, for applications where the stability of the phosphorylated derivative is critical, a serum-stable derivative will be used. Similar considerations apply to modified Hu-IFN- β . For applications where stability in serum is not essential, the serum-unstable phosphorylated -- or analog -- may be used.

The applications described herein are not limited to proteins phosphorylated at the serine residue; it has been described above how kinases phosphorylate other amino acids such as threonine or tyrosine (20, 21, 23). Thus, proteins modified at these amino acids are within the contemplation of the invention. Because of the configuration of such derivatized labelled proteins, it is not to be excluded that their stability in serum may be improved if the corresponding serine-phosphorylated derivative is not adequately serum-stable.

2. General Diagnostic Reagents.

Additional specific applications of the modified proteins of the invention are noteworthy. As referred to herein, virtually all proteins can be engineered to introduce single or multiple phosphorylation (or analog) sites. Such proteins can be used for a wide variety of scientific purposes: to study the fate of these proteins in animals or humans; to study their stabilities; or for use as any laboratory reagent where a radioactive protein is useful.

For example, molecular weight standards are commonly used for polyacrylamide gel electrophoresis. Proteins with phosphorylation sites would make convenient autoradiographic markers such as molecular weight markers, isoelectric focusing markers or other markers. For such applications the serum stability is generally not critical, nor is the retention of the biological activity of the protein, e.g., the interferon. Thus, for certain uses or applications it is not essential that a phosphorylatable protein in accordance with the invention have biological activity.

3. Anticancer Therapeutic "Bomb".

A particularly noteworthy and interesting application made possible by the invention is what has been called here in the vernacular, a therapeutic or more specifically an antitumor "therapeutic radiation bomb". Such a biologically-active composition uses biotin coupled to a tumor-specific monoclonal antibody (Mab) (or to Fab or Fab' fragments if more appropriate), and a multiple "modified" streptavidin bound to each Mab-bound biotin, each streptavidin being modified in that it has multiple phosphorylated groups. Since streptavidin is itself a tetramer, multiple radioactive groups are thus provided. These multiple radioactive groups expose the tumor with radiation which is greatly amplified and hence more readily detectable and would produce greater tumor destruction. In the case where it is highly phosphorylatable it is much more easily detectable. Thus, each one of the biotins which is bound to each tumor-specific Mab binds tightly to

the multiple streptavidin molecules which in turn contain multiple labelled phosphorus atoms, or their equivalent isotopes.

It is evident that depending on the therapeutic or diagnostic objectives, all streptavidins may be radioactive-phosphorus labelled or partially or totally radioactive-thiophosphorus labelled, or labelled with different phosphorus or sulfur isotopes, which have different decay modes or levels of radiation energy. Such isotopes are discussed below.

Because antibody molecules are themselves multichain molecules, many sites can be introduced into the antibodies or Fab fragments directly by the procedures of this invention.

4. Hormones, Cytokines, Lymphokines, Growth Factors.

Hormones labelled with radioactive phosphorus or sulfur are another class of biological materials within the scope of this invention. For instance, phosphorylated (e.g., ^{33}P , ^{32}P) hormones can be bound to specific cell types differentially over other tissues. Cancerous tissues containing increased number of receptors for such hormones can be treated with appropriately phosphorylated hormones which will then specifically bind to these cells; thus therapy will be significantly improved.

Further, labelled hormones are commonly used for receptor studies to examine their binding to cell surface receptors, to soluble receptors or other reagents and materials.

Typical of the labelled hormones (^{33}P , ^{32}P) contemplated by the invention are growth hormone, insulin, FSH, LH, and others. It is evident such hormones genetically constructed lend themselves to the introduction of one or more putative phosphorylatable or thiophosphorylatable groups.

As noted above for hormones, the same considerations apply to cytokines, lymphokines, growth factors (i.e., IL-1, IL-2, IL-3, TNF- α , TNF- β , the various CSF molecules, erythropoietin EGF, NGF and others) and any proteins with cell and/or tissue specificity to one degree or another.

5. Antibodies.

Streptavidin labelled by means of phosphorylation may be used directly to enhance immunoassays as a substitute for unlabelled streptavidin or enzyme-linked unlabelled streptavidin. The invention also contemplates introducing phosphorus or analog labels into genetically engineered antibodies (see references 93-99), more particularly Mabs, or in the Fab or Fab' fragment. Such Mabs are useful for diagnostic and therapeutic purposes. The phosphorylated Mabs can be made to target specific tumor-associated antigens or a variety of tumors, like breast and colon cancer cells, malignant melanoma cells, ovarian carcinoma cells, and other malignant tumors.

6. Further Therapeutic Uses.

Other uses contemplated in accordance with the invention are as follows: Monoclonal or appropriate cocktails of antibodies and/or antibody fragments (such as the Fab or Fab' fragments) are fruitful molecules in which in accordance with the invention phosphorylation or other labellable sites can be introduced. The use of ^{32}P in therapy has been demonstrated for polycythemia vera and other malignancies (116). Thus, it is clear that the high energy β particle is effective as an anticellular agent. The attachment of ^{32}P through the introduction of phosphorylation site(s) in Mabs or their appropriate fragments (Fab and Fab') would also be effective for the therapy of tumors to which these monoclonal antibodies are specific. A large number of monoclonal antibodies have been developed to tumor-associated antigens from breast, colon, ovarian, and other adenocarcinomas, malignant melanoma, and many other tumors. Thus, Mabs directed to the tumor associated antigens of these tumors are expected to be highly effective when labelled with ^{32}P . The labelling can be increased by use of cassettes of phosphorylation sites or directly by introduction of multiple phosphorylation sites into the intact protein or the appropriate fragments through genetic engineering. By "cassette" is meant a multifunctional moiety.

When multiple labelled phosphorylation sites are introduced in accordance with the invention in Mabs, this may reduce the binding specificity and/or affinity of the modified Mabs for the specific epitope targeted. It can be seen that under such circumstances the use of a biotinylated Mab linked to the multiple phosphorylated streptavidin (as described above) has distinct advantages; the specificity of the Mab is not altered and yet the radioactivity of the diagnostic agent has been many-fold enhanced.

The invention also has implications for the preparation of therapeutic agents to which patients are likely to develop an adverse antigenic response. Thus, the monoclonal antibodies can be engineered successively in accordance with the invention with different phosphorylation sites. When introduced into patients who have become sensitive to or who are producing antibodies to the injected antibody because of the phosphorylation site, then by changing to a different phosphorylation site, the antigenic character of the protein can be modified. Thus, it may be possible to use such antibodies in multiple successive therapeutic regimens in patients who are reacting with the antibody of the previous type. For this purpose a series of antibodies with a variety of phosphorylation sites can be developed. Each series would be

designed to have a different epitopic structure and be used sequentially. Alternatively a cocktail of such different antibodies can be used initially so that any one is present at a fraction of the total. This would minimize antibody formation to any one of the new sites.

7. Various Isotopes.

In accordance with the invention, as discussed above, phosphorylated derivatives should be serum-stable for certain applications. Various isotopes can be employed that are more effective than others for a specific therapeutic purpose. For example, ^{33}P may be substituted for ^{32}P in the phosphorylation reaction. It is less likely that ^{35}S with a half-life of about 87 days would be normally as useful as an anticellular reagent because it is a low energy β emitter. Nevertheless, conceivably there may be specific uses for ^{35}S labelled monoclonal antibodies in therapy and/or diagnosis.

Table 3 below shows various isotopes (and other pertinent particulars) which are especially useful for introduction into proteins in accordance with the invention.

Table 3

Isotopes for Labellable Groups			
Isotope	Half-Life	Type of Decay	Energy of Radiation
^{32}P	14.2 days	β^-	1.707 Mev
^{33}P	24.4 days	β^-	0.25 Mev
^{35}S	87.0 days	β^-	0.167 Mev
^{38}S	2.87 hours	β^-	1.1 Mev

Decay factors and radioactivity at any given time is available in the literature. For instance, for comparison between ^{125}I and ^{131}I with, on the other hand, ^{32}P and ^{35}S , reference is made to Appendices, Table A.1.7 (Supplement 2) in Current Protocols in Molecular Biology, cited (43).

Thus, the invention provides tailored-designed proteins for specific biological purposes.

An important implication of this invention is the greater safety of the labelled Mabs due to lower energy emission levels and the nature of the radio emission. Specifically, Mabs labelled with ^{32}P or ^{33}P have significantly lower energy emission levels than conventional radio-labels for protein such as ^{125}I ; moreover, the decay emission of the phosphorus and sulfur isotopes (^{32}P , ^{33}P , ^{35}S and ^{38}S) is beta particles, as compared to gamma rays of ^{125}I as are common in existing labelling protocols.

The safety feature of the beta-labelled proteins, e.g., Mabs or streptavidins (as discussed) in accordance with the invention, is very significant for diagnostic and therapeutic uses of the invention. Beta emitters penetrate the tumor but are not emitted as readily as gamma ray emitters from the patient to surrounding medical staff and non-medical attending individuals.

By selecting ^{35}S (which has a half-life of 87 days) and the ^{35}S phosphate ATP analog to ^{32}P one can significantly increase the effective radioactive life of the therapeutic agent.

Thus, the proteins labelled in accordance with the invention have a spectrum of meaningful advantageous properties heretofore not readily available.

The invention is not limited to the use of unstable isotopes. In the future it may be advantageous to label a protein with a stable isotope that would be suitable for detection by NMR, nuclear activation, or future developed procedures. Nor is it necessary that the label be a "radio" label providing it is an identifiable label.

8. Radioimmunoassays with Labelled Antigens.

In accordance with the invention the phosphorylated proteins can be generally used as the radio-labelled component. These radioimmunoassays can be used with polyclonal as well as with monoclonal antibodies. If the introduction of a new phosphorylation site into a protein changes the antigenic structure of the protein in the area of the phosphorylation site, or even at distant linear positions of the protein, and alters the antigenic behavior, the protein in accordance with the invention, can be modified to introduce a phosphorylation site at a different position so that the antigenic behavior will remain stable and for the protein to bind with the polyclonal or monoclonal antibody of interest.

Thus, the invention provides considerable versatility regarding the position where the label can be introduced. Generally it will be preferred to introduce the phosphorus (or other radio-label) at a site that will not disrupt the antigen-antibody binding.

9. Sandwich Radioimmunoassays.

In sandwich radioimmunoassays with monoclonal antibodies, the introduction of phosphorylation sites into an antibody in accordance with the invention is a sensitive method to follow the binding of the second antibody. Thus, the sensitivity of such sandwich radioimmunoassays can be increased substantially. Particularly, when multiple phosphorylation sites are introduced in accordance with the invention into the protein directly or by the addition of a fusion phosphorylation cassette, the sensitivity of such assays will be increased many-fold.

Another advantage of the invention is to be noted. Because the phosphorylation reaction is gentle, unlike the iodination or other chemical modifications necessary to radio-label proteins with iodine or other reagents, monoclonal antibodies that are inactivated by the chemical or iodination procedures are not likely to be inactivated by the phosphorylation procedure. Thus, the process of the invention allows for the phosphorylation of proteins normally too sensitive for labelling with iodine. The introduction of a phosphate analog with ^{35}S provides a radio-labelled protein derivative with a long half-life (1.5 times longer than ^{125}I and 6 times longer than ^{32}P). Thus, when Mabs are labelled with ^{35}S , they will have a substantially longer shelf-life compared to the ^{32}P or ^{125}I radio-labelled derivatives.

As discussed above, the invention allows for the selection of the most appropriate labelling isotope, as compared to ^{125}I , for instance.

10. Imaging.

Generally for imaging of tumors or tissues in an animal or a patient, a high energy gamma emitter is generally preferable to a high energy β emitter, which by and large would be absorbed by the tissues. However, in certain imaging studies in animals or in patients, Mabs to which ^{32}P , ^{33}P or ^{35}S are attached through introduced phosphorylation sites in accordance with the invention may be useful.

For example, it can be seen that Mabs labelled with ^{32}P , ^{33}P or ^{35}S could be useful in *in vivo* studies in which biopsy specimens are to be examined. The spread of a tumor during surgery could be followed by utilizing a radioisotope detector probe to follow the local spread of the tumor and guide the extent of the surgery. In addition, tissue specimens which are fixed or frozen can be taken to which these proteins will remain bound (that is, antibodies to the tumor-associated antigens or other ligands). Thus, autoradiographs of tissue sections can provide information about the extent of tumor spread and the extent of binding of specific monoclonal antibodies to tumor-associated antigens can be thoroughly evaluated. Furthermore, as an *in vitro* reagent with cells or tissue slices, such labelled antibodies would be highly sensitive reagents to detect tumor-associated antigens or other antigens by the usual types of assays employed.

11. Anti-antibodies.

There are many known uses for anti-antibodies such as anti-mouse, anti-human, anti-sheep, and anti-goat antibodies, etc. or monoclonal antibodies as single entities or as a cocktail. Such antibodies can be engineered in accordance with the invention to introduce single or multiple phosphorylation sites and, accordingly labelled with a variety of isotopes as described above. These provide general reagents where anti-antibodies are necessary, particularly in radioimmunoassays, autoradiography, or any other reactions in which anti-antibodies are useful.

12. Rapid Purification of Phosphorylated Proteins.

The invention has also applications in separating and purifying proteins. Proteins which are phosphorylated can be separated from those which are not; proteins which are more phosphorylated than others can be separated.

For instance, where proteins can be phosphorylated, it is common for only a percentage of the molecules to be phosphorylated. The total phosphorylation, of course, can be enhanced by the introduction of multiple phosphorylation sites in the protein in accordance with the invention so that few molecules escape phosphorylation. To be able to separate the phosphorylated from the non-phosphorylated proteins is especially useful for molecules with a single phosphorylation site where there may be phosphorylated and non-phosphorylated molecules in the population. In this manner, the effectiveness of any phosphorylated derivatives is increased. Separation of phosphorylated from non-phosphorylated molecules can be accomplished by developing polyclonal or monoclonal antibodies to the phosphorylation sites with and/or without derivatized phosphate groups. Such polyclonal and monoclonal antibodies are expected to have considerable value in purifying the proteins and have been described (see for example 119-124).

13. Dephosphorylation of Proteins.

Considerable emphasis has been placed herein on aspects of phosphorylation. It is a consequence of the phosphorylation (with phosphate or thiophosphate groups) that the removal of the label is also facilitated in that dephospho-

rylation is a milder procedure which tends to be less disruptive of the protein molecule than procedures in the prior art for removal of ^{125}I from proteins. Thus, in cases where it is useful to remove the radioisotope, this can be achieved relatively easily and gently by an enzyme reaction. A variety of phosphatases can be used for this purpose. Most phosphatases have comparatively low specificity (for example, reference 100, pages 192-193, 203, 223-224, 736-739) although a few have very high specificity such as those acting on sugar phosphates and the enzyme that dephosphorylates glycogen synthetase b and phosphorylase b (47, 100; also reference 101 pages 372-373, for example). Furthermore, specific dephosphorylation of phosphorylated proteins can be achieved by reversal of the reaction of protein-serine and -tyrosine kinases (107). If it is necessary to determine whether in fact the phosphate addition causes a change in the activity of the protein, rather than aging, denaturation, or other manipulations, the phosphate can be removed and the activity of the protein again determined. In such a manner, a definitive understanding of the effect of phosphorylation on the activity of the protein can be assessed. This may be useful in determining the activities of various phosphorylated interferons.

The concept of "dephosphorylation" has an interesting application which is essentially the "converse" of that taught herein. Wherever a site in a protein in the native state is naturally phosphorylatable the removal of that site would be particularly desirable when it is known that the naturally phosphorylatable protein causes some undesired results. An illustration would be proteins associated with oncogenic viruses such as Rous sarcoma virus (RSV) and cellular oncogenes.

14. Phosphorylation Cassettes.

The invention also contemplates an alternative method for labelling proteins without inserting the coding sequence for the phosphorylation site (or cassette) into the nucleotide coding sequence of the protein, and yet still use the invention. This procedure would be particularly useful for large proteins like immunoglobulins for use in various assays. Such alternative method calls for a polypeptide which is phosphorylated to be chemically linked to the large protein. The linking would be by any bifunctional reagent or an activated derivative (like N-hydroxysuccinimide), as is known in the art.

This technique could use a polypeptide with multiple phosphorylation sites in tandem or "cassette" that can be introduced within or at either end of a protein. The DNA coding for the tandem phosphorylation sites would be flanked by restriction sites for easy cleaving and insertion into the DNA containing the coding sequence for the protein to be linked to the larger protein. Such a phosphorylation cassette could be expressed as a small polypeptide then phosphorylated and then chemically linked to the larger protein.

15. Phosphorylatable Human or Animal Donor Genes.

Further, it is within the contemplation of the invention to provide DNA sequences engineered into appropriate vectors or cell lines or even into animals by transgenic techniques. Thus cells or animals could produce phosphorylatable (and/or phosphorylated) proteins such as immunoglobulins after phosphorylation sites are introduced into the proteins by the methods of this invention. Phosphorylatable chimeric antibodies with a mouse variable region and human constant region could be developed (93-99). The human antibodies used as the donor molecule would be engineered to contain single or multiple phosphorylation sites. Analogously, this could be applied to proteins other than immunoglobulins.

16. Use of Phosphorylation Sites to Map Tertiary Structure of Proteins.

By introducing a small phosphorylation recognition site into a protein randomly along the entire linear protein chain, it will be possible to obtain information about the tertiary structure of proteins. The sequence encoding the phosphorylation site is inserted randomly within the DNA sequence encoding the protein of interest. The insertion must be made in such a way that the phosphorylation sequence is in phase with upstream and downstream codons so that an insertion for a phosphorylation site is made without interrupting the phase of translation. The expressed protein, therefore, contains the identical linear sequence of the original protein with an insertion of a phosphorylation site in a given position along the chain. By generating a large series of insertions (ideally after every amino acid position of the protein chain), it is possible to determine whether the kinase recognizes the sequence in the context in which it is placed by a simple assay to determine the rate and extent of phosphorylation at that position. The rate and extent of phosphorylation depends on the accessibility of that site to the phosphokinase which reflects its position in the tertiary structure (outside, internal, buried, etc.). A complete linear map of the accessibility of the phosphorylation sites along the entire chain will provide an outline of the structural features of the protein that are inside and outside in the tertiary structural configuration. Insertions of amino acids should be designed to minimize perturbations. In some cases insertion of a phosphorylation site can occur by simply changing one or more amino acids rather than inserting several amino acids comprising the phosphorylation recognition site.

The generation of such DNA insertions to make the appropriate variety of insertional mutant proteins can be done

in many ways. Insertions can be introduced along a protein chain systematically or randomly by methods comparable to saturation mutagenesis. Alternatively, rather than generating mutants from a given DNA sequence by inserting the sequence encoding the phosphorylation site into the DNA, one can generate synthetic oligonucleotides so that the entire DNA chain is synthesized *de novo*. Combinations of these procedures and general cloning strategies could easily provide an entire bank of new mutant proteins with phosphorylation sites distributed linearly along the chain.

This procedure provides information about the tertiary structure and folding of the protein in solution. It compliments methods such as x-ray crystallography which will provide tertiary structure information of the proteins in the crystal. Furthermore, the method will be useful to determine the tertiary structure of proteins which have resisted efforts to obtain appropriate crystals for determination of X-ray crystallographic structures.

This aspect is an illustration of a protein having numerous putative phosphorylation sites, ideally after each amino acid in the sequence of the protein; and the corresponding phosphorylated protein. Likewise, this is an illustration of a DNA sequence encoding the putative phosphorylation site(s) inserted in the DNA sequence encoding the selected protein of interest.

17. Other Applications.

There are other applications for the labelled proteins of the invention. In general virtually any protein that contains a label (radio-label, fluorescent-label, chemical-label, enzyme-label, etc.) can alternatively be labelled with phosphate by the introduction of phosphorylation site(s) in accordance with the invention. The purification of such proteins can be followed in a sensitive assay by simply measuring the ability to accept a phosphate group rather than to follow enzyme activity. Such proteins engineered in accordance with the invention, therefore, can be purified easily and themselves be used as a tracer to follow the purification of other proteins to which they are similar. For example, it is likely that a protein with a single phosphorylation site engineered with very little modification of the protein structure itself would be purified similarly to the unmodified protein.

In practice, by having a stock of phosphorylatable proteins or series of markers, the labelled derivatives can be prepared conveniently by the simple phosphorylation reaction when desired. Thus, the proteins of the invention which are phosphorylatable provide a useful inventory of the corresponding labelled proteins.

18. Pharmaceutical and Biologically Active Compositions.

The modified proteins of the invention can be formulated according to known methods to prepare pharmaceutically useful compositions. For instance, the human alpha interferon-like protein hereof is combined in admixture with a pharmaceutically acceptable carrier vehicle. Suitable vehicles and their formulation are described in Remington's Pharmaceutical Sciences by E. W. Martin, which is hereby incorporated by reference. Such compositions will contain an effective amount of the interferon-like protein or other proteins hereof together with a suitable amount of vehicle in order to prepare pharmaceutically acceptable compositions suitable for effective administration to the host. The host can be a mammal or not. The carrier may be liquid, solid, or other. Of course therapeutic applications for humans and veterinary applications are intended for the biologically active compositions of the invention. The biologically active composition of the invention is to be administered in a biologically or therapeutically effective amount which can be readily determined by one skilled in the art. Generally it is the smallest amount for which a desired response will be obtained to an amount which is excessive for practical or other purposes.

The biologically active compositions of the invention can also include any other biologically active substance which does not adversely affect the desired activity, particularly the activity or use of the modified protein of the invention.

It is understood that the modified proteins of the invention can be obtained by chemical and/or enzymatic synthesis rather than by recombinant DNA technology.

While reference has been made to particular preferred embodiments and to several uses and applications made possible by the invention, it will be understood that the present invention is not to be construed as limited to such, but rather to the lawful scope of the appended claims and subject matter covered by the doctrine of equivalents.

From the description provided hereinabove it will be appreciated by one skilled in the art that the invention makes a significant and meritorious contribution to the art.

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Claims

Claims for the following Contracting States : DE, FR, GB, IT, NL, SE, CH, BE, LU, AT

1. A protein which is modified to comprise an amino acid sequence recognizable by a protein kinase, which amino acid sequence is, as part of the protein, phosphorylatable and was not present, or was unavailable for phosphorylation, in the unmodified protein.
2. The phosphorylatable protein of claim 1 wherein the said modification is provided by recombinant DNA techniques.

3. The phosphorylatable protein of claim 2 which is a functional protein.
4. The phosphorylatable protein of claim 2 which is an interferon-like protein selected from the group consisting of an interferon-like Hu-IFN- α protein or an interferon-like Hu-IFN- β protein.
5. The phosphorylatable Hu-IFN- α -like protein of claim 4.
6. The phosphorylatable Hu-IFN- β -like protein of claim 4.
7. The phosphorylatable Hu-IFN- α of claim 4 wherein the interferon-like Hu-IFN- α protein is a fused Hu-IFN- α /gamma protein.
8. The phosphorylatable fused Hu-IFN- β -like protein of claim 7 which has the amino acid sequence shown schematically in FIG. 3.
9. The phosphorylatable Hu-IFN- α -like protein of claim 5 which is selected from the group of the three amino acid sequences shown in FIG. 8 and identified as α A-P1, α A-P2, and α A-P3.
10. The phosphorylatable Hu-IFN- α of claim 7 wherein the interferon-like Hu-IFN- α protein possesses a biological property of Hu-IFN- α and/or of Hu-IFN-gamma.
11. The phosphorylatable Hu-IFN- α -like protein of claim 4 which possesses a biological property of Hu-IFN- α and/or of Hu-IFN- β .
12. The phosphorylatable Hu-IFN- α -like protein or Hu-IFN- β -like protein of claim 4 which comprises an amino acid consensus sequence which is recognizable by protein kinase.
13. The phosphorylatable Hu-IFN- α -like protein of claim 12 which comprises an amino acid consensus sequence which is an insert into a phosphorylatable Hu-IFN- α -like protein.
14. The phosphorylatable Hu-IFN- α -like protein of claim 12 wherein the amino acid consensus sequence recognizable by the protein kinase is fused to the carboxyl terminus of the amino acid sequence of Hu-IFN- α .
15. The phosphorylatable interferon-like Hu-IFN- α protein of claim 12 which contains at least two amino acid segments, a first segment which is the amino acid sequence of natural Hu-IFN- α , and a second amino acid segment which contains the amino acid consensus sequence which is recognizable by protein kinase, which second segment may be incorporated with in or linked to the first amino and segment.
16. The phosphorylatable Hu-IFN- α -like protein of claim 5 which is phosphorylated and contains a serum stable phosphate.
17. The phosphorylatable Hu-IFN- α -like protein of claim 5 which upon phosphorylation is stable in human serum.
18. The phosphorylatable Hu-IFN- α -like protein of claim 12 wherein the amino acid consensus sequence contains Ser.
19. The phosphorylatable Hu-IFN- α -like protein of claim 12 wherein the amino acid consensus sequence contains an amino acid selected from the group consisting of Ser, Thr or Tyr.
20. The phosphorylatable Hu-IFN- α -like protein of claim 12 wherein the amino acid consensus sequence is Arg-Arg-Ala-Ser-.
21. The phosphorylatable Hu-IFN- α -like protein of claim 12 wherein the amino acid consensus sequence contains Arg-Arg-Ala-Ser-Xaa, wherein Xaa is selected from the group of Gln, Val or Leu.
22. The phosphorylatable Hu-IFN- α -like protein of claim 12 wherein the protein kinase is the cAMP-dependent protein kinase.
23. The phosphorylatable Hu-IFN- α -like protein of claim 4 which is purified.

24. The phosphorylatable Hu-IFN- α -like protein of claim 4 which is phosphorylated with a radioactive label.
25. The phosphorylatable Hu-IFN- α -like protein of claim 24 wherein the label is a beta ray-emitting atom.
- 5 26. The phosphorylatable Hu-IFN- α -like protein of claim 25 wherein the atom is selected from the group consisting of phosphorus and sulfur.
27. The phosphorylatable Hu-IFN- α -like protein of claim 26 wherein the radioactive label is ^{32}P or ^{33}P .
- 10 28. A phosphorylatable Hu-IFN- α -like protein or a Hu-IFN- β -like protein, the phosphorylated protein being defined in any one of claims 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21 or 23.
29. The phosphorylatable Hu-IFN- α -like protein having the amino acid sequence shown in FIG. 3 or one of the amino acid sequences identified as $\alpha\text{A-P1}$, $\alpha\text{A-P2}$, and $\alpha\text{A-P3}$, shown in FIG. 8.
- 15 30. The phosphorylatable Hu-IFN- β -like protein of the phosphorylated protein defined in claim 6.
31. A recombinant DNA sequence which encodes a protein which has been modified to contain an amino acid consensus sequence which is recognizable by protein kinase, which protein is not phosphorylatable absent said modification.
- 20 32. The recombinant DNA sequence of claim 31 which contains a DNA sequence which encodes a functional protein which is normally not phosphorylatable and a DNA sequence which encodes the amino acid consensus sequence which is recognizable by protein kinase and which contains the putative labelling site.
- 25 33. The recombinant DNA sequence of claim 32 wherein the functional protein is an interferon-like protein selected from the group consisting of Hu-IFN- α or Hu-IFN- β .
34. The recombinant DNA sequence of claim 32 wherein the second nucleotide sequence has one codon which is common with the first nucleotide sequence.
- 30 35. A recombinant DNA sequence which is selected from the group consisting of the DNA sequences encoding a protein selected from the group consisting of a Hu-IFN- α -like protein or a Hu-IFN- β -like protein, which protein has been modified to contain an amino acid consensus sequence recognisable by protein kinase comprising a putative phosphorylatable site.
- 35 36. The DNA recombinant sequence of claim 35 which encodes a phosphorylatable Hu-IFN- α -like protein.
37. A recombinant DNA cloning vector useful for expressing an exogenous functional protein which vector includes the DNA sequence which encodes a Hu-IFN- α -like protein or a Hu-IFN- β -like protein which protein has been modified to contain an amino acid consensus sequence recognisable by protein kinase comprising a putative phosphorylation site, the phosphorylatable protein being an interferon-like Hu-IFN- α protein or an interferon-like Hu-IFN- β protein.
- 40 38. The cloning vector of claim 37 wherein the encoded protein is the interferon-like protein Hu-IFN- α .
- 45 39. The cloning vector of claim 37 wherein the encoded protein is the interferon-like protein Hu-IFN- β .
40. The cloning vector of claim 37 which contains the DNA sequence coding for a fused Hu-IFN- α/γ protein.
41. The cloning vector of claim 37 which contains the DNA sequence coding for the modified interferons referred to as -P1, -P2, and -P3 in FIG. 8, respectively.
- 50 42. A transformed host cell capable of expressing a functional protein which host cell contains a recombinant DNA vector encoding a Hu-IFN- α -like protein or a Hu-IFN- β -like protein, which has been modified to contain an amino acid consensus sequence recognisable by protein kinase.
- 55 43. A biologically active composition which comprises a biologically compatible carrier and in a biologically effective amount, a phosphorylated protein defined in any one of claims 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26 or 27.

44. The phosphorylatable protein of claim 1 which possesses a biological property of the unphosphorylatable protein.
45. The phosphorylatable protein of claim 1 which contains at least two amino acid segments, a first segment which is the amino acid sequence of the unphosphorylatable protein, and a second amino acid segment which contains the amino acid sequence recognizable by the protein kinase, which second segment may be incorporated within or linked to the first amino acid segment.
46. The phosphorylatable protein of claim 1 which is phosphorylated and contains a serum stable phosphate.
47. The protein of claim 1 which is a phosphorylatable monoclonal antibody.
48. The monoclonal antibody of claim 47 which is a chimeric, a hybrid or a modified monoclonal antibody.
49. The monoclonal antibody of claim 47 which is tumor-specific.
50. The monoclonal antibody of claim 47 wherein the site for phosphorylation is in the Fab or Fab¹ fragment thereof.
51. The monoclonal antibody of any one of claims 47 to 50 which is phosphorylated.
52. The monoclonal antibody of any one of claims 47 to 51 which is radiolabelled.
53. The protein of claim 1 which is a cytokine.
54. The protein of claim 1 which is tumor necrosis factor (TNF).
55. The protein of claim 54 which is selected from the group consisting of TNF- α and TNF- β .
56. A method for producing a phosphorylatable protein from an otherwise unphosphorylatable protein, comprising the steps of:
 - identifying an amino acid sequence recognizable by a protein kinase (hereafter "PK");
 - defining and constructing a nucleotide sequence which codes for said amino acid PK recognition sequence;
 - determining a nucleotide sequence coding for said unphosphorylatable protein;
 - constructing a modified nucleotide sequence encoding said phosphorylatable protein;
 - expressing said modified nucleotide sequence to produce said phosphorylatable protein.
57. The method of claim 56 wherein said step of constructing said modified nucleotide sequence includes affixing said nucleotide sequence encoding said amino acid PK recognition sequence to said nucleotide sequence encoding said unphosphorylatable protein.
58. The method of claim 56 wherein said step of constructing said modified nucleotide sequence includes introducing said nucleotide sequence encoding said amino acid PK recognition sequence in said nucleotide sequence encoding said unphosphorylatable protein.
59. The method of claim 56 wherein said amino acid PK recognition sequence identified in said identifying step contains Ser.
60. The method of claim 56 wherein said amino acid PK recognition sequence identified in said identifying step contains an amino acid selected from the group consisting of Ser, Thr or Tyr.
61. The method of claim 56 wherein said amino acid PK recognition sequence identified in said identifying step is Arg-Arg-Ala-Ser-.
62. The method of claim 56 wherein said amino acid PK recognition sequence identified in said identifying step contains Arg-Arg-Ala-Ser-Xaa, wherein Xaa is selected from the group of all amino acids.
63. The method of claim 56 wherein the protein kinase of said identifying step is the cAMP-dependent protein kinase.

64. The method of claim 56 wherein said phosphorylatable protein is phosphorylated with a radioactive label.

Claims for the following Contracting State : ES

1. A method for producing a protein which is modified to comprise an amino acid sequence recognizable by a protein kinase, which amino acid sequence is, as part of the protein, phosphorylatable and was not present, or was unavailable for phosphorylation, in the unmodified protein.
2. The method of claim 1 wherein the said modification is provided by recombinant DNA techniques.
3. The method of claim 2 which is a functional protein.
4. The method of claim 2 which is an interferon-like protein selected from the group consisting of an interferon-like Hu-IFN- α protein or an interferon-like Hu-IFN- β protein.
5. The method of Claim 4 wherein the protein is Hu-IFN- α -like.
6. The method of Claim 4 wherein the protein is Hu-IFN- β -like.
7. The method of claim 4 wherein the interferon-like Hu-IFN- α protein is a fused Hu-IFN- α /gamma protein.
8. The method of Claim 7 wherein the Hu-IFN- β -like protein has the amino acid sequence shown schematically in FIG. 3.
9. The method of Claim 7 wherein the Hu-IFN- α -like protein is selected from the group of the three amino acid sequences shown in FIG. 8 and identified as α A-P1, α A-P2, and α A-P3.
10. The method of claim 7 wherein the interferon-like Hu-IFN- α protein possesses a biological property of Hu-IFN- α and/or of Hu-IFN-gamma.
11. The method of Claim 4 wherein the Hu-IFN- α -like protein possesses a biological property of Hu-IFN- α and/or of Hu-IFN- β .
12. The method of Claim 4 wherein the Hu-IFN- α -like protein or Hu-IFN- β -like protein comprises an amino acid consensus sequence which is recognizable by protein kinase.
13. The method of Claim 12 wherein the Hu-IFN- α -like protein comprises an amino acid consensus sequence which is an insert into a phosphorylatable Hu-IFN- α -like protein.
14. The method of Claim 12 wherein the Hu-IFN- α -like protein wherein the amino acid consensus sequence recognizable by the protein kinase is fused to the carboxyl terminus of the amino acid sequence of Hu-IFN- α .
15. The method of Claim 12 wherein the interferon-like Hu-IFN- α protein contains at least two amino acid segments, a first segment which is the amino acid sequence of natural Hu-IFN- α , and a second amino acid segment which contains the amino acid consensus sequence which is recognizable by protein kinase, which second segment may be incorporated within or linked to the first amino acid segment.
16. The method of Claim 5 wherein the Hu-IFN- α -like protein is phosphorylated and contains a serum stable phosphate.
17. The method of Claim 5 wherein the Hu-IFN- α -like protein upon phosphorylation is stable in human serum.
18. The method of Claim 12 wherein the amino acid consensus sequence of the Hu-IFN- α -like protein contains Ser.
19. The method of Claim 12 wherein the amino acid consensus sequence of the Hu-IFN- α -like protein contains an amino acid selected from the group consisting of Ser, Thr or Tyr.
20. The method of Claim 12 wherein the amino acid consensus sequence of the Hu-IFN- α -like protein is

Arg-Arg-Ala-Ser-

21. The method of Claim 12 wherein the amino acid consensus sequence of the Hu-IFN- α -like protein contains Arg-Arg-Ala-Ser-Xaa, wherein Xaa is selected from the group of Gln, Val or Leu.

22. The method of Claim 12 wherein the protein is Hu-IFN- α -like protein and the protein kinase is the cAMP-dependent protein kinase.

23. The method of Claim 4 further comprising purifying the phosphorylatable Hu-IFN- α -like protein.

24. The method of claim 4 further comprising phosphorylating phosphorylated Hu-IFN- α -like protein with a radioactive label.

25. The method of claim 24 wherein the label is a beta ray-emitting atom.

26. The method of claim 25 wherein the atom is selected from the group consisting of phosphorus and sulfur.

27. The method of claim 26 wherein the radioactive label is ^{32}P or ^{33}P .

28. A method of producing a Hu-IFN- α -like protein or a Hu-IFN- β -like protein, the phosphorylated protein being defined in any one of claims 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21 or 23, the method comprising the step of (a) introducing into a protein which is not phosphorylatable an amino acid sequence recognizable by a protein kinase for phosphorylation or (b) synthesising the said phosphorylatable protein from amino.

29. A method of producing The phosphorylatable Hu-IFN- α -like protein having the amino acid sequence shown in FIG. 3 or one of the amino acid sequences identified as $\alpha\text{A-P1}$, $\alpha\text{A-P2}$, and $\alpha\text{A-P3}$, shown in FIG. 8, the method comprising the step of (a) introducing into a protein which is not phosphorylatable an amino acid sequence recognizable by a protein kinase for phosphorylation or (b) synthesising the said phosphorylatable protein from amino.

30. A method of producing The phosphorylatable Hu-IFN- β -like protein of the phosphorylated protein defined in claim 6, the method comprising the step of (a) introducing into a protein which is not phosphorylatable an amino acid sequence recognizable by a protein kinase for phosphorylation or (b) synthesising the said phosphorylatable protein from amino.

31. A method of producing a recombinant DNA sequence which encodes a modified protein as defined in Claim 1 by recombinant DNA techniques.

32. The method of Claim 31 wherein the recombinant DNA sequence contains a DNA sequence which encodes a functional protein which is normally not phosphorylatable and a DNA sequence which encodes the amino acid consensus sequence which is recognizable by protein kinase and which contains the putative labelling site.

33. The method of Claim 32 wherein the functional protein is an interferon-like protein selected from the group consisting of Hu-IFN- α or Hu-IFN- β .

34. The method of Claim 32 wherein the second nucleotide sequence has one codon which is common with the first nucleotide sequence.

35. A method for producing a recombinant DNA sequence which is selected from the group consisting of the DNA sequences encoding a protein selected from the group consisting of a Hu-IFN- α -like protein or a Hu-IFN- β -like protein, which protein has been modified to contain an amino acid consensus sequence recognisable by protein kinase comprising a putative phosphorylatable site by recombinant DNA techniques.

36. The method according to Claim 35 wherein DNA recombinant sequence encodes a phosphorylatable Hu-IFN- α -like protein.

37. A method of producing a recombinant DNA cloning vector useful for expressing an exogenous functional protein which vector includes the DNA sequence which encodes a Hu-IFN- α -like protein or a Hu-IFN- β -like protein which protein has been modified to contain an amino acid consensus sequence recognisable by protein kinase comprising

a putative phosphorylation site, the phosphorylatable protein being an interferon-like Hu-IFN- α protein or an interferon-like Hu-IFN- β protein by recombinant DNA techniques.

38. The method of claim 37 wherein the encoded protein is the interferon-like protein Hu-IFN- α .
39. The method of claim 37 wherein the encoded protein is the interferon-like protein Hu-IFN- β .
40. The method of claim 37 wherein the cloning vector contains the DNA sequence coding for a fused Hu-IFN- α /gamma protein.
41. The method of claim 37 wherein the cloning vector contains the DNA sequence coding for the modified interferons referred to as -P1, -P2, and -P3 in FIG. 8, respectively.
42. A method for producing a transformed host cell capable of expressing a functional protein which host cell contains a recombinant DNA vector encoding a Hu-IFN- α -like protein or a Hu-IFN- β -like protein, which has been modified to contain an amino acid consensus sequence recognisable by protein kinase comprising transforming the host cell with the recombinant DNA vector.
43. A method of making A biologically active composition comprising the steps of admixing a biologically compatible carrier and a biologically effective amount, a phosphorylated protein as defined in any one of claims 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26 or 27.
44. A method according to claim 1 wherein The phosphorylatable protein possesses a biological property of the unphosphorylatable protein.
45. A method according to claim 1 wherein The phosphorylatable protein contains at least two amino acid segments, a first segment which is the amino acid sequence of the unphosphorylatable protein, and a second amino acid segment which contains the amino acid sequence recognizable by the protein kinase, which second segment may be incorporated within or linked to the first amino acid segment.
46. A method according to claim 1 wherein The phosphorylatable protein is phosphorylated and contains a serum stable phosphate.
47. A method according to claim 1 wherein The protein is a phosphorylatable monoclonal antibody.
48. A method according to claim 47 wherein The monoclonal antibody is a chimeric, a hybrid or a modified monoclonal antibody.
49. A method according to Claim 47 wherein The monoclonal antibody is tumor-specific.
50. A method according to Claim 47 wherein the site for phosphorylation is in the Fab or Fab¹ fragment thereof.
51. A method according to any one of claims 47 to 50 further comprising phosphorylating the monocloned antibody.
52. A method according to any one of claims 47 to 51 further comprising radiolabelling the monoclonal antibody.
53. A method according to Claim 1 wherein the protein is a cytokine.
54. A method according to Claim 1 wherein the protein is tumor necrosis factor (TNF) or interleukin -3 (IL-3).
55. A method according to Claim 54 wherein the protein is selected from the group consisting of TNF- α and TNF- β .
56. A method for producing a phosphorylatable protein from an otherwise unphosphorylatable protein, comprising the steps of:

identifying an amino acid sequence recognizable by a protein kinase (hereafter "PK");
defining and constructing a nucleotide sequence which codes for said amino acid PK recognition sequence;
determining a nucleotide sequence coding for said unphosphorylatable protein

constructing a modified nucleotide sequence encoding said phosphorylatable protein;
expressing said modified nucleotide sequence to produce said phosphorylatable protein.

57. The method of claim 56 wherein said step of constructing said modified nucleotide sequence includes affixing said nucleotide sequence encoding said amino and PK recognition sequence to said nucleotide sequence encoding said unphosphorylatable protein.
58. The method of claim 56 wherein said step of constructing said modified nucleotide sequence includes introducing said nucleotide sequence encoding said amino acid PK recognition sequence in said nucleotide sequence encoding said unphosphorylatable protein.
59. The method of claim 56 wherein said amino acid PK recognition sequence identified in said identifying step contains Ser.
60. The method of claim 56 wherein said amino acid PK recognition sequence identified in said identifying step contains an amino acid selected from the group consisting of Ser, Thr or Tyr.
61. The method of claim 56 wherein said amino acid PK recognition sequence identified in said identifying step is Arg-Arg-Ala-Ser-.
62. The method of claim 56 wherein said amino acid PK recognition sequence identified in said identifying step contains Arg-Arg-Ala-Ser-Xaa, wherein Xaa is selected from the group of all amino acids.
63. The method of claim 56 wherein the protein kinase of said identifying step is the cAMP-dependent protein kinase.
64. The method of claim 56 wherein said phosphorylatable protein is phosphorylated with a radioactive label.

Claims for the following Contracting State : GR

1. A method for producing a protein which is modified to comprise an amino acid sequence recognizable by a protein kinase, which amino acid sequence is, as part of the protein, phosphorylatable and was not present, or was unavailable for phosphorylation, in the unmodified protein.
2. The method of claim 1 wherein the said modification is provided by recombinant DNA techniques.
3. The method of claim 2 which is a functional protein.
4. The method of claim 2 which is an interferon-like protein selected from the group consisting of an interferon-like Hu-IFN- α protein or an interferon-like Hu-IFN- β protein.
5. The method of Claim 4 wherein the protein is Hu-IFN- α -like.
6. The method of Claim 4 wherein the protein is Hu-IFN- β -like.
7. The method of claim 4 wherein the interferon-like Hu-IFN- α protein is a fused Hu-IFN- α /gamma protein.
8. The method of Claim 7 wherein the Hu-IFN- β -like protein has the amino acid sequence shown schematically in FIG. 3.
9. The method of Claim 7 wherein the Hu-IFN- α -like protein is selected from the group of the three amino acid sequences shown in FIG. 8 and identified as α A-P1, α A-P2, and α A-P3.
10. The method of claim 7 wherein the interferon-like Hu-IFN- α protein possesses a biological property of Hu-IFN- α and/or of Hu-IFN-gamma.
11. The method of Claim 4 wherein the Hu-IFN- α -like protein possesses a biological property of Hu-IFN- α and/or of Hu-IFN- β .

12. The method of Claim 4 wherein the Hu-IFN- α -like protein or Hu-IFN- β -like protein comprises an amino acid consensus sequence which is recognizable by protein kinase.
13. The method of Claim 12 wherein the Hu-IFN- α -like protein comprises an amino acid consensus sequence which is an insert into a phosphorylatable Hu-IFN- α -like protein.
14. The method of Claim 12 wherein the Hu-IFN- α -like protein wherein the amino acid consensus sequence recognizable by the protein kinase is fused to the carboxyl terminus of the amino acid sequence of Hu-IFN- α .
15. The method of Claim 12 wherein the interferon-like Hu-IFN- α protein contains at least two amino acid segments, a first segment which is the amino acid sequence of natural Hu-IFN- α , and a second amino acid segment which contains the amino acid consensus sequence which is recognizable by protein kinase, which second segment may be incorporated within or linked to the first amino acid segment.
16. The method of Claim 5 wherein the Hu-IFN- α -like protein is phosphorylated and contains a serum stable phosphate.
17. The method of Claim 5 wherein the Hu-IFN- α -like protein upon phosphorylation is stable in human serum.
18. The method of Claim 12 wherein the amino acid consensus sequence of the Hu-IFN- α -like protein contains Ser.
19. The method of Claim 12 wherein the amino acid consensus sequence of the Hu-IFN- α -like protein contains an amino acid selected from the group consisting of Ser, Thr or Tyr.
20. The method of Claim 12 wherein the amino acid consensus sequence of the Hu-IFN- α -like protein is Arg-Arg-Ala-Ser-.
21. The method of Claim 12 wherein the amino acid consensus sequence of the Hu-IFN- α -like protein contains Arg-Arg-Ala-Ser-Xaa, wherein Xaa is selected from the group of Gln, Val or Leu.
22. The method of Claim 12 wherein the protein is Hu-IFN- α -like protein and the protein kinase is the cAMP-dependent protein kinase.
23. The method of Claim 4 further comprising purifying the phosphorylatable Hu-IFN- α -like protein.
24. The method of claim 4 further comprising phosphorylating phosphorylated Hu-IFN- α -like protein with a radioactive label.
25. The method of claim 24 wherein the label is a beta ray-emitting atom.
26. The method of claim 25 wherein the atom is selected from the group consisting of phosphorus and sulfur.
27. The method of claim 26 wherein the radioactive label is ^{32}P or ^{33}P .
28. A method of producing a Hu-IFN- α -like protein or a Hu-IFN- β -like protein, the phosphorylated protein being defined in any one of claims 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21 or 23, the method comprising the step of (a) introducing into a protein which is not phosphorylatable an amino acid sequence recognizable by a protein kinase for phosphorylation or (b) synthesising the said phosphorylatable protein from amino acid precursors.
29. A method of producing The phosphorylatable Hu-IFN- α -like protein having the amino acid sequence shown in FIG. 3 or one of the amino acid sequences identified as $\alpha\text{A-P1}$, $\alpha\text{A-P2}$, and $\alpha\text{A-P3}$, shown in FIG. 8, the method comprising the step of (a) introducing into a protein which is not phosphorylatable an amino acid sequence recognizable by a protein kinase for phosphorylation or (b) synthesising the said phosphorylatable protein from amino acid precursors.
30. A method of producing The phosphorylatable Hu-IFN- β -like protein of the phosphorylated protein defined in claim 6, the method comprising the step of (a) introducing into a protein which is not phosphorylatable an amino acid sequence recognizable by a protein kinase for phosphorylation or (b) synthesising the said phosphorylatable protein from amino acid precursors.

31. A recombinant DNA sequence which encodes a modified protein as defined in Claim 1.
32. The recombinant DNA sequence of claim 31 which contains a DNA sequence which encodes a functional protein which is normally not phosphorylatable and a DNA sequence which encodes the amino acid consensus sequence which is recognizable by protein kinase and which contains the putative labelling site.
33. The recombinant DNA sequence of claim 32 wherein the functional protein is an interferon-like protein selected from the group consisting of Hu-IFN- α or Hu-IFN- β .
34. The recombinant DNA sequence of claim 32 wherein the second nucleotide sequence has one codon which is common with the first nucleotide sequence.
35. A recombinant DNA sequence which is selected from the group consisting of the DNA sequences encoding a protein selected from the group consisting of a Hu-IFN- α -like protein or a Hu-IFN- β -like protein, which protein has been modified to contain an amino acid consensus sequence recognisable by protein kinase comprising a putative phosphorylatable site.
36. The DNA recombinant sequence of claim 35 which encodes a phosphorylatable Hu-IFN- α -like protein.
37. A recombinant DNA cloning vector useful for expressing an exogenous functional protein which vector includes the DNA sequence which encodes a Hu-IFN- α -like protein or a Hu-IFN- β -like protein which protein has been modified to contain an amino acid consensus sequence recognisable by protein kinase comprising a putative phosphorylation site, the phosphorylatable protein being an interferon-like Hu-IFN- α protein or an interferon-like Hu-IFN- β protein.
38. The cloning vector of claim 37 wherein the encoded protein is the interferon-like protein Hu-IFN- α .
39. The cloning vector of claim 37 wherein the encoded protein is the interferon-like protein Hu-IFN- β .
40. The cloning vector of claim 37 which contains the DNA sequence coding for a fused Hu-IFN- α /gamma protein.
41. The cloning vector of claim 37 which contains the DNA sequence coding for the modified interferons referred to as -P1, -P2, and -P3 in FIG. 8, respectively.
42. A transformed host cell capable of expressing a functional protein which host cell contains a recombinant DNA vector encoding a Hu-IFN- α -like protein or a Hu-IFN- β -like protein which has been modified to contain an amino acid consensus sequence recognisable by protein kinase.
43. A method of making A biologically active composition comprising the steps of admixing a biologically compatible carrier and a biologically effective amount, a phosphorylated protein as defined in any one of claims 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26 or 27.
44. A method according to claim 1 wherein The phosphorylatable protein possesses a biological property of the unphosphorylatable protein.
45. A method according to claim 1 wherein The phosphorylatable protein contains at least two amino acid segments, a first segment which is the amino acid sequence of the unphosphorylatable protein, and a second amino acid segment which contains the amino acid sequence recognizable by the protein kinase, which second segment may be incorporated within or linked to the first amino acid segment.
46. A method according to Claim 1 wherein The phosphorylatable protein is phosphorylated and contains a serum stable phosphate.
47. A method according to Claim 1 wherein The protein is a phosphorylatable monoclonal antibody.
48. A method according to Claim 47 wherein The monoclonal antibody is a chimeric, a hybrid or a modified monoclonal antibody.
49. A method according to Claim 47 wherein The monoclonal antibody is tumor-specific.

50. A method according to Claim 47 wherein the site for phosphorylation is in the Fab or Fab¹ fragment thereof.
51. A method according to any one of claims 47 to 50 further comprising phosphorylating the monoclonal antibody.
52. A method according to any one of claims 47 to 51 further comprising radiolabelling the monoclonal antibody.
53. A method according to Claim 1 wherein the protein is a cytokine.
54. A method according to Claim 1 wherein the protein is tumor necrosis factor (TNF) or interleukin-3 (IL-3).
55. A method according to Claim 54 wherein the protein is selected from the group consisting of TNF- α and TNF- β .
56. A method for producing a phosphorylatable protein from an otherwise unphosphorylatable protein, comprising the steps of:
- identifying an amino acid sequence recognizable by a protein kinase (hereafter "PK");
 defining and constructing a nucleotide sequence which codes for said amino acid PK recognition sequence;
 determining a nucleotide sequence coding for said unphosphorylatable protein;
 constructing a modified nucleotide sequence encoding said phosphorylatable protein;
 expressing said modified nucleotide sequence to produce said phosphorylatable protein.
57. The method of claim 56 wherein said step of constructing said modified nucleotide sequence includes affixing said nucleotide sequence encoding said amino acid PK recognition sequence to said nucleotide sequence encoding said unphosphorylatable protein.
58. The method of claim 56 wherein said step of constructing said modified nucleotide sequence includes introducing said nucleotide sequence encoding said amino acid PK recognition sequence in said nucleotide sequence encoding said unphosphorylatable protein.
59. The method of claim 56 wherein said amino acid PK recognition sequence identified in said identifying step contains Ser.
60. The method of claim 56 wherein said amino acid PK recognition sequence identified in said identifying step contains an amino acid selected from the group consisting of Ser, Thr or Tyr.
61. The method of claim 56 wherein said amino acid PK recognition sequence identified in said identifying step is Arg-Arg-Ala-Ser-.
62. The method of claim 56 wherein said amino acid PK recognition sequence identified in said identifying step contains Arg-Arg-Ala-Ser-Xaa, wherein Xaa is selected from the group of all amino acids.
63. The method of claim 56 wherein the protein kinase of said identifying step is the cAMP-dependent protein kinase.
64. The method of claim 56 wherein said phosphorylatable protein is phosphorylated with a radioactive label.

Patentansprüche

Patentansprüche für folgende Vertragsstaaten : DE, FR, GB, IT, NL, SE, CH, BE, LU, AT

1. Protein, welches derart modifiziert ist, daß es eine durch eine Proteinkinase erkennbare, als Teil des Proteins phosphorylierbare und in dem nicht modifizierten Protein nicht vorhandene oder für die Phosphorylierung nicht verfügbare Aminosäuresequenz umfaßt.
2. Phosphorylierbares Protein nach Anspruch 1, wobei für die Modifikation durch rekombinante DNA-Techniken gesorgt wurde.

3. Phosphorylierbares Protein nach Anspruch 2, bei welchem es sich um ein funktionelles Protein handelt.
4. Phosphorylierbares Protein nach Anspruch 2, bei dem es sich um ein interferonartiges Protein, ausgewählt aus der Gruppe ein interferonartiges Hu-IFN- α Protein oder ein interferonartiges Hu-IFN- β Protein handelt.
5. Phosphorylierbares Hu-IFN- α -artiges Protein nach Anspruch 4.
6. Phosphorylierbares Hu-IFN- β -artiges Protein nach Anspruch 4.
7. Phosphorylierbares Hu-IFN- α Protein nach Anspruch 4, wobei es sich bei dem interferonartigen HU-IFN- α Protein um ein fusioniertes Hu-IFN- α /gamma Protein handelt.
8. Phosphorylierbares fusioniertes Hu-IFN- β -artiges Protein nach Anspruch 7, welches die schematisch in Figur 3 dargestellte Aminosäuresequenz aufweist.
9. Phosphorylierbares Hu-IFN- α -artiges Protein nach Anspruch 5, welches aus der Gruppe der drei in Figur 8 dargestellten Aminosäuresequenzen ausgewählt und als α A-P1, α A-P2 und α A-P3 identifiziert ist.
10. Phosphorylierbares Hu-IFN- α . Protein nach Anspruch 7, wobei das interferonartige Hu-IFN- α protein eine biologische Eigenschaft von Hu-IFN- α und/oder von Hu-IFN-gamma aufweist.
11. Phosphorylierbares Hu-IFN- α -artiges Protein nach Anspruch 4, welches eine biologische Eigenschaft von Hu-IFN- α und/oder von Hu-IFN- β aufweist.
12. Phosphorylierbares Hu-IFN- α -artiges Protein oder Hu-IFN- β -artiges Protein nach Anspruch 4, welches eine durch eine Proteinkinase erkennbare Aminosäurekonsenssequenz umfaßt.
13. Phosphorylierbares Hu-IFN- α -artiges Protein nach Anspruch 12, welches eine ein Insert in ein phosphorylierbares Hu-IFN- α -artiges Protein darstellende Aminosäurekonsenssequenz umfaßt.
14. Phosphorylierbares Hu-IFN- α -artiges Protein nach Anspruch 12, wobei die durch die Proteinkinase erkennbare Aminosäurekonsenssequenz mit dem Carboxylende der Aminosäuresequenz von Hu-IFN- α fusioniert ist.
15. Phosphorylierbares interferonartiges Hu-IFN- α Protein nach Anspruch 12, welches mindestens zwei Aminosäuresegmente, nämlich ein erstes Segment, bei dem es sich um die Aminosäuresequenz von natürlichem Hu-IFN- α handelt, und ein zweites Aminosäuresegment, das die durch Proteinkinase erkennbare Aminosäurekonsenssequenz enthält, aufweist, wobei das zweite Segment in das erste Aminosäuresegment eingebaut oder mit diesem verbunden sein kann.
16. Phosphorylierbares Hu-IFN- α -artiges Protein nach Anspruch 5, welches phosphoryliert ist und ein serumstables Phosphat enthält.
17. Phosphorylierbares Hu-IFN- α -artiges Protein nach Anspruch 5, welches nach der Phosphorylierung in Humanserum stabil ist.
18. Phosphorylierbares Hu-IFN- α -artiges Protein nach Anspruch 12, wobei die Aminosäurekonsenssequenz Ser enthält.
19. Phosphorylierbares Hu-IFN- α -artiges Protein nach Anspruch 12, wobei die Aminosäurekonsenssequenz eine Aminosäure, ausgewählt aus der Gruppe Ser, Thr oder Tyr, enthält.
20. Phosphorylierbares Hu-IFN- α -artiges Protein nach Anspruch 12, wobei die Aminosäurekonsenssequenz Arg-Arg-Ala-Ser- ist.
21. Phosphorylierbares Hu-IFN- α -artiges Protein nach Anspruch 12, wobei die Aminosäurekonsenssequenz Arq-Arg-Ala-Ser-Xaa mit Xaa ausgewählt aus der Gruppe Gln, Val oder Leu enthält.
22. Phosphorylierbares Hu-IFN- α -artiges Protein nach Anspruch 12, wobei es sich bei der Proteinkinase um die

cAMP-abhängige Proteinkinase handelt.

23. Phosphorylierbares Hu-IFN- α -artiges Protein nach Anspruch 4 in gereinigter Form.
- 5 24. Phosphorylierbares Hu-IFN- α -artiges Protein nach Anspruch 4, das mit einer radioaktiven Markierung phosphoryliert ist.
25. Phosphorylierbares Hu-IFN- α -artiges Protein nach Anspruch 24, wobei es sich bei der Markierung um ein β -Strahlung emittierendes Atom handelt.
- 10 26. Phosphorylierbares Hu-IFN- α -artiges Protein nach Anspruch 25, wobei das Atom aus der Gruppe Phosphor und Schwefel ausgewählt ist.
- 15 27. Phosphorylierbares Hu-IFN- α -artiges Protein nach Anspruch 26, wobei es sich bei der radioaktiven Markierung um ^{32}P oder ^{33}P handelt.
28. Phosphorylierbares Hu-IFN- α -artiges Protein oder Hu-IFN- β -artiges Protein, wobei das phosphorylierte Protein der Definition gemäß einem der Ansprüche 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21 oder 23 entspricht.
- 20 29. Phosphorylierbares Hu-IFN- α -artiges Protein mit der in Figur 3 dargestellten Aminosäuresequenz oder einer der als A-P1, α A-P2 und α A-P3 identifizierten und in Figur 8 dargestellten Aminosäuresequenzen.
30. Phosphorylierbares Hu-IFN- β -artiges Protein des phosphorylierten Proteins gemäß Anspruch 6.
- 25 31. Rekombinante DNA-Sequenz mit Kodierung für ein modifiziertes Protein nach Anspruch 1.
32. Rekombinante DNA-Sequenz nach Anspruch 31, enthaltend eine DNA-Sequenz mit Kodierung für ein funktionelles Protein, welches normalerweise nicht phosphorylierbar ist, und eine DNA-Sequenz mit Kodierung für die Aminosäurekonsenssequenz, die durch eine Proteinkinase erkennbar ist und die mutmaßliche Markierungsstelle enthält.
- 30 33. Rekombinante DNA-Sequenz nach Anspruch 32, wobei es sich bei dem funktionellen Protein um ein interferonartiges Protein aus der Gruppe Hu-IFN- α oder Hu-IFN- β handelt.
- 35 34. Rekombinante DNA-Sequenz nach Anspruch 32, wobei die Zweite Nukleotidsequenz ein mit der ersten Nukleotidsequenz gemeinsames Kodon aufweist.
- 40 35. Rekombinante DNA-Sequenz, ausgewählt aus der Gruppe DNA-Sequenzen mit Kodierung für ein Protein, ausgewählt aus der Gruppe Hu-IFN- α -artiges Protein oder Hu-IFN- β -artiges Protein, wobei das Protein derart modifiziert wurde, daß es eine durch eine Proteinkinase erkennbare Aminosäurekonsenssequenz mit einer mutmaßlichen phosphorylierbaren Stelle enthält.
36. Rekombinante DNA-Sequenz nach Anspruch 35 mit Kodierung für ein phosphorylierbares Hu-IFN- α -artiges Protein.
- 45 37. Rekombinanter DNA-Klonierungsvektor mit der Eignung zum Expressieren eines exogenen funktionellen Proteins, wobei der Vektor die DNA-Sequenz mit Kodierung für ein Hu-IFN- α -artiges Protein oder ein Hu-IFN- β -artiges Protein beinhaltet, das Protein derart modifiziert wurde, daß es eine durch eine Proteinkinase erkennbare Aminosäurekonsenssequenz mit einer mutmaßlichen Phosphorylierungsstelle enthält und das phosphorylierbare Protein aus einem interferonartigen Hu-IFN- α Protein oder einem interferonartigen Hu-IFN- β Protein besteht.
- 50 38. Klonierungsvektor nach Anspruch 37, wobei es sich bei dem kodierten Protein um das interferonartige Protein Hu-IFN- α handelt.
39. Klonierungsvektor nach Anspruch 37, wobei es sich bei dem kodierten Protein um das interferonartige Protein Hu-IFN- β handelt.
- 55 40. Klonierungsvektor nach Anspruch 37, enthaltend die DNA-Sequenz mit Kodierung für ein fusioniertes Hu-IFN- α /gamma Protein.

41. Klonierungsvektor nach Anspruch 37, enthaltend die DNA-Sequenz mit Kodierung für die in Figur 8 mit -P1, -P2 und bzw. -P3 bezeichneten modifizierten Interferone.

42. Transformierte Wirtszelle mit der Fähigkeit zum Expressieren des funktionellen Proteins, wobei die Wirtszelle einen rekombinanten DNA-Vektor mit Kodierung für ein Hu-IFN- α -artiges Protein oder ein Hu-IFN- β -artiges Protein, welches derart modifiziert wurde, daß es eine durch eine Proteinkinase erkennbare Aminosäurekonsensussequenz enthält, enthält.

43. Biologisch aktive Zusammensetzung, umfassend einen biologisch verträglichen Träger und - in biologisch wirksamer Menge - ein phosphoryliertes Protein nach einem der Ansprüche 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26 oder 27.

44. Phosphorylierbares Protein nach Anspruch 1, welches eine biologische Eigenschaft des nicht phosphorylierbaren Proteins aufweist.

45. Phosphorylierbares Protein nach Anspruch 1, enthaltend mindestens zwei Aminosäuresegmente, nämlich ein erstes Segment, bei dem es sich um die Aminosäuresequenz des nicht phosphorylierbaren Proteins handelt, und ein zweites Aminosäuresegment, welches die durch die Proteinkinase erkennbare Aminosäuresequenz enthält, wobei das zweite Segment in das erste Aminosäuresegment eingebaut oder mit diesem verbunden sein kann.

46. Phosphorylierbares Protein nach Anspruch 3, welches phosphoryliert ist und ein serumstables Phosphat enthält.

47. Phosphorylierbares Protein nach Anspruch 1, bei dem es sich um einen phosphorylierbaren monoklonalen Antikörper handelt.

48. Monoklonaler Antikörper nach Anspruch 47, bei welchem es sich um eine Chimäre, einen Hybrid oder modifizierten monoklonalen Antikörper handelt.

49. Monoklonaler Antikörper nach Anspruch 47, welcher Tumorspezifisch ist.

50. Monoklonaler Antikörper nach Anspruch 47, wobei die Phosphorylierungsstelle in dem Fab- oder dem Fab¹-Fragment hiervon enthalten ist.

51. Monoklonaler Antikörper nach einem der Ansprüche 47 bis 50, welcher phosphoryliert ist.

52. Monoklonaler Antikörper nach einem der Ansprüche 47 bis 51, welcher radioaktiv markiert ist.

53. Protein nach Anspruch 1, bei dem es sich um ein Zytokin handelt.

54. Protein nach Anspruch 1, bei welchem es sich um den Tumornekrose-Faktor (TNF) oder Interleukin-3 (IL-3) handelt.

55. Protein nach Anspruch 54, ausgewählt aus der Gruppe TNF- α und TNF- β .

56. Verfahren zur Herstellung eines phosphorylierbaren Proteins aus einem ansonst nicht phosphorylierbaren Protein, umfassend folgende Stufen:

Identifizieren einer durch eine Proteinkinase (im folgenden als "PK" bezeichnet) erkennbaren Aminosäuresequenz;

Definieren und konstruieren einer Nukleotidsequenz mit kodierung für die Aminosäure-PK-Erkennungssequenz;

Bestimmen einer Nukleotidsequenz mit Kodierung für das nicht phosphorylierbare Protein;

Konstruieren einer modifizierten Nukleotidsequenz mit Kodierung für das phosphorylierbare Protein und

Expressieren der modifizierten Nukleotidsequenz zur Produktion des phosphorylierbaren Proteins.

57. Verfahren nach Anspruch 56, wobei die Stufe der Konstruktion der modifizierten Nukleotidsequenz ein Anhängen der Nukleotidsequenz mit Kodierung für die Aminosäure-PK-Erkennungssequenz an die Nukleotidsequenz mit Kodierung für das nicht phosphorylierbare Protein beinhaltet.
- 5 58. Verfahren nach Anspruch 56, wobei die Stufe der Konstruktion der modifizierten Nukleotidsequenz eine Einfügung der Nukleotidsequenz mit Kodierung für die Aminosäure-PK-Erkennungssequenz in die Nukleotidsequenz mit Kodierung für das nicht phosphorylierbare Protein beinhaltet.
- 10 59. Verfahren nach Anspruch 56, wobei die in der Identifizierungsstufe identifizierte Aminosäure-PK-Erkennungssequenz Ser enthält.
60. Verfahren nach Anspruch 56, wobei die in der Identifizierungsstufe identifizierte Aminosäure-PK-Erkennungssequenz eine Aminosäure, ausgewählt aus der Gruppe Ser, Thr oder Tyr enthält.
- 15 61. Verfahren nach Anspruch 56, wobei die in der Identifizierungsstufe identifizierte Aminosäure-PK-Erkennungssequenz aus Arg-Arg-Ala-Ser- besteht.
62. Verfahren nach Anspruch 56, wobei die in der Identifizierungsstufe identifizieren Aminosäure-PK-Erkennungssequenz Arg-Arg-Ala-Ser-Xaa, mit Xaa ausgewählt aus der Gruppe sämtlicher Aminosäuren, enthält.
- 20 63. Verfahren nach Anspruch 56, wobei es sich bei der Proteinkinase der Identifizierungsstufe um die cAMP-abhängige Proteinkinase handelt.
- 25 64. Verfahren nach Anspruch 56, wobei das phosphorylierbare Protein mit einer radioaktiven Markierung phosphoryliert ist.

Patentansprüche für folgenden Vertragsstaat : ES

- 30 1. Verfahren zur Herstellung eines Proteins, welches derart modifiziert ist, daß es eine durch eine Proteinkinase erkennbare, als Teil des Proteins phosphorylierbare und in dem nicht modifizierten Protein nicht vorhandene oder für die Phosphorylierung nicht verfügbare Aminosäuresequenz umfaßt.
- 35 2. Verfahren nach Anspruch 1, wobei für die Modifikation durch rekombinante DNA-Techniken gesorgt wurde.
3. Verfahren nach Anspruch 2, wobei es sich bei dem Protein um ein funktionelles Protein handelt.
4. Verfahren nach Anspruch 2, wobei es sich bei dem Protein um ein interferonartiges Protein, ausgewählt aus der Gruppe ein interferonartiges Hu-IFN- α Protein oder ein interferonartiges Hu-IFN- β Protein handelt.
- 40 5. Verfahren nach Anspruch 4, wobei das Protein Hu-IFN- α -artig ist.
6. Verfahren nach Anspruch 4, wobei das Protein Hu-IFN- β -artig ist.
- 45 7. Verfahren nach Anspruch 4, wobei es sich bei dem interferonartigen Hu-IFN- α Protein um ein fusioniertes Hu-IFN- α /gamma Protein handelt.
8. Verfahren nach Anspruch 7, wobei das Hu-IFN- β -artige Protein die schematisch in Figur 3 dargestellte Aminosäuresequenz aufweist.
- 50 9. Verfahren nach Anspruch 7, wobei das Hu-IFN- α -artige Protein aus der Gruppe der in Figur 8 dargestellten und als α A-P1, α A-P2 und α A-P3 identifizierten drei Aminosäuresequenzen ausgewählt ist.
- 55 10. Verfahren nach Anspruch 7, wobei das interferonartige Hu-IFN- α protein eine biologische Eigenschaft von Hu-IFN- α und/oder von Hu-IFN-gamma aufweist.
11. Verfahren nach Anspruch 4, wobei das Hu-IFN- α -artige Protein eine biologische Eigenschaft von Hu-IFN- α und/oder

von Hu-IFN- β aufweist.

12. Verfahren nach Anspruch 4, wobei das Hu-IFN- α -artige Protein oder Hu-IFN- β -artige Protein eine durch eine Proteinkinase erkennbare Aminosäurekonsenssequenz umfaßt.

13. Verfahren nach Anspruch 12, wobei das Hu-IFN- α -artige Protein eine ein Insert in ein phosphorylierbares Hu-IFN- α -artiges Protein darstellende Aminosäurekonsenssequenz umfaßt.

14. Verfahren nach Anspruch 12, wobei bei dem Hu-IFN- α -artigen protein die durch die Proteinkinase erkennbare Aminosäurekonsenssequenz mit dem Carboxylende der Aminosäuresequenz von Hu-IFN- α fusioniert ist.

15. Verfahren nach Anspruch 12, wobei das interferonartige Hu-IFN- α Protein mindestens zwei Aminosäuresegmente, nämlich ein erstes segment, bei dem es sich um die Aminosäuresequenz von natürlichem Hu-IFN- α handelt, und ein zweites Aminosäuresegment, das die durch Proteinkinase erkennbare Aminosäurekonsenssequenz enthält, aufweist, und wobei das zweite Segment in das erste Aminosäuresegment eingebaut oder mit diesem verbunden sein kann.

16. Verfahren nach Anspruch 5, wobei das Hu-IFN- α -artige Protein phosphoryliert ist und ein serumstables Phosphat enthält.

17. Verfahren nach Anspruch 5, wobei das Hu-IFN- α -artige Protein nach der Phosphorylierung in Humanserum stabil ist.

18. Verfahren nach Anspruch 12, wobei die Aminosäurekonsenssequenz des Hu-IFN- α -artigen Proteins Ser enthält.

19. Verfahren nach Anspruch 12, wobei die Aminosäurekonsenssequenz des Hu-IFN- α -artigen Proteins eine Aminosäure, ausgewählt aus der Gruppe Ser, Thr oder Tyr, enthält.

20. Verfahren nach Anspruch 12, wobei die Aminosäurekonsenssequenz des Hu-IFN- α -artigen Proteins aus Arg-Arg-Ala-Ser- besteht.

21. Verfahren nach Anspruch 12, wobei die Aminosäurekonsenssequenz des Hu-IFN- α -artigen Proteins Arg-Arg-Ala-Ser-Xaa mit Xaa ausgewählt aus der Gruppe Gln, Val oder Leu enthält.

22. Verfahren nach Anspruch 12, wobei das Protein aus einem Hu-IFN- α -artigen Protein und die Proteinkinase aus der cAMP-abhängigen Proteinkinase bestehen.

23. Verfahren nach Anspruch 4, weiterhin umfassend die Reinigung des phosphorylierbaren Hu-IFN- α -artigen Proteins.

24. Verfahren nach Anspruch 4, weiter umfassend die Phosphorylierung des phosphorylierbaren Hu-IFN- α -artigen Proteins mit einer radioaktiven Markierung.

25. Verfahren nach Anspruch 24, wobei es sich bei der Markierung um ein β -Strahlung emittierendes Atom handelt.

26. Verfahren nach Anspruch 25, wobei das Atom aus der Gruppe Phosphor und Schwefel ausgewählt ist.

27. Verfahren nach Anspruch 26, wobei es sich bei der radioaktiven Markierung um ^{32}P oder ^{33}P handelt.

28. Verfahren zur Herstellung eines Hu-IFN- α -artigen Proteins oder eines Hu-IFN- β -artigen Proteins, wobei das phosphorylierte Protein der Definition gemäß einem der Ansprüche 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21 oder 23 entspricht und das Verfahren folgende Stufen umfaßt:

(a) Einführen einer durch eine Proteinkinase zur Phosphorylierung erkennbaren Aminosäuresequenz in ein nicht phosphorylierbares Protein oder

(b) Synthese des phosphorylierbaren Proteins aus Aminosäurevorläufern.

29. Verfahren zur Herstellung des phosphorylierbaren Hu-IFN- α -artigen Proteins mit der in Figur 3 dargestellten Ami-

nosäuresequenz oder einer der als α A-P1, α A-P2 und α A-P3 identifizierten und in Figur 8 dargestellten Aminosäuresequenzen, wobei das Verfahren folgende Stufen umfaßt:

- 5 (a) Einführen einer durch eine Proteinkinase zur Phosphorylierung erkennbaren Aminosäuresequenz in ein nicht phosphorylierbares Protein oder
 - (b) Synthese des phosphorylierbaren Proteins aus Aminosäurevorläufern.
- 10 **30.** Verfahren zur Herstellung eines phosphorylierbaren Hu-IFN- β -artigen Proteins des phosphorylierten Proteins gemäß Anspruch 6 wobei das Verfahren folgende Stufen umfaßt:
- (a) Einführen einer durch eine Proteinkinase zur Phosphorylierung erkennbaren Aminosäuresequenz in ein nicht phosphorylierbares Protein oder
 - 15 (b) Synthese des phosphorylierbaren Proteins aus Aminosäurevorläufern.
- 31.** Verfahren zur Herstellung einer rekombinanten DNA-Sequenz mit Kodierung für ein modifiziertes Protein gemäß der Definition von Anspruch 1 durch rekombinante DNA-Technik.
- 20 **32.** Verfahren nach Anspruch 31, enthaltend eine DNA-Sequenz mit Kodierung für ein funktionelles Protein, welches normalerweise nicht phosphorylierbar ist, und eine DNA-Sequenz mit Kodierung für die Aminosäurekonsenssequenz, die durch eine Proteinkinase erkennbar ist und die mutmaßliche Markierungsstelle enthält.
- 25 **33.** Verfahren nach Anspruch 32, wobei es sich bei dem funktionellen Protein um ein interferonartiges Protein, ausgewählt aus der Gruppe Hu-IFN- α oder Hu-IFN- β handelt.
- 34.** Verfahren nach Anspruch 32, wobei die zweite Nukleotidsequenz ein mit der ersten Nukleotidsequenz gemeinsames Kodon aufweist.
- 30 **35.** Verfahren zur Herstellung einer rekombinanten DNA-Sequenz, ausgewählt aus der Gruppe DNA-Sequenzen mit Kodierung für ein Protein, ausgewählt aus der Gruppe Hu-IFN- α -artiges Protein oder Hu-IFN- β -artiges Protein, wobei das Protein Tierart modifiziert wurde, daß es eine durch eine Proteinkinase erkennbare Aminosäurekonsenssequenz mit einer mutmaßlichen phosphorylierbaren Stelle enthält, durch rekombinante DNA-Techniken.
- 35 **36.** Verfahren nach Anspruch 35, wobei die rekombinante DNA-Sequenz für ein phosphorylierbares Hu-IFN- α -artiges Protein kodiert.
- 37.** Verfahren zur Herstellung eines rekombinanten DNA-Klonierungsvektors mit der Eignung zum Exprimieren eines exogenen funktionellen Proteins, wobei der Vektor die DNA-Sequenz mit Kodierung für ein Hu-IFN- α -artiges Protein oder ein Hu-IFN- β -artiges Protein beinhaltet, das Protein derart modifiziert wurde, daß es eine durch eine Proteinkinase erkennbare Aminosäurekonsenssequenz mit einer mutmaßlichen Phosphorylierungsstelle enthält und das phosphorylierbare Protein aus einem interferonartigen Hu-IFN- α Protein oder einem interferonartigen Hu-IFN- β Protein besteht, durch rekombinante DNA-Techniken.
- 40 **38.** Verfahren nach Anspruch 37, wobei es sich bei dem kodierten Protein um das interferonartige Protein Hu-IFN- α handelt.
- 39.** Verfahren nach Anspruch 37, wobei es sich bei dem kodierten Protein um das interferonartige Protein Hu-IFN- β handelt.
- 50 **40.** Verfahren nach Anspruch 37, wobei der Klonierungsvektor die DNA-Sequenz mit Kodierung für ein fusioniertes Hu-IFN- α A/gamma Protein enthält.
- 41.** Verfahren nach Anspruch 37, wobei der Klonierungsvektor die DNA-Sequenz mit Kodierung für die in Figur 8 mit -P1, -P2 und bzw. -P3 bezeichneten modifizierten Interferone enthält.
- 55 **42.** Verfahren zur Herstellung einer transformierten Wirtszelle mit der Fähigkeit zum Exprimieren des funktionellen Proteins, wobei die Wirtszelle einen rekombinanten DNA-Vektor mit Kodierung für ein Hu-IFN- α -artiges Protein oder ein

Hu-IFN- β -artiges Protein, welches derart modifiziert wurde, daß es eine durch eine Proteinkinase erkennbare Aminosäurekonsensussequenz enthält, aufweist, umfassend die Transformation der Wirtszelle mit dem rekombinanten DNA-Vektor.

- 5 **43.** Verfahren zur Zubereitung einer biologisch aktiven Zusammensetzung, umfassend die Stufe des Vermischen eines biologisch verträglichen Trägers und einer biologisch wirksamen Menge eines phosphorylierten Proteins entsprechend der Definition gemäß einem der Ansprüche 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26 oder 27.
- 10 **44.** Verfahren nach Anspruch 1, wobei das phosphorylierbare Protein eine biologische Eigenschaft des nicht phosphorylierbaren Proteins aufweist.
- 15 **45.** Verfahren nach Anspruch 1, wobei das phosphorylierbare Protein mindestens zwei Aminosäuresegmente, nämlich ein erstes Segment, bei dem es sich um die Aminosäuresequenz des nicht phosphorylierbaren Proteins handelt, und ein zweites Aminosäuresegment, welches die durch die Proteinkinase erkennbare Aminosäuresequenz enthält, wobei das zweite Segment in das erste Aminosäuresegment eingebaut oder mit diesem verbunden sein kann.
- 20 **46.** Verfahren nach Anspruch 1, wobei das phosphorylierbare Protein phosphoryliert ist und ein serumstabiles Phosphat enthält.
- 25 **47.** Verfahren nach Anspruch 1, wobei das Protein aus einem phosphorylierbaren monoklonalen Antikörper besteht.
- 30 **48.** Verfahren nach Anspruch 47, wobei der monoklonale Antikörper aus einer Chimäre, einem Hybrid oder modifizierten monoklonalen Antikörper besteht.
- 35 **49.** Verfahren nach Anspruch 47, wobei der monoklonale Antikörper Tumor-spezifisch ist.
- 40 **50.** Verfahren nach Anspruch 47, wobei die Phosphorylierungsstelle in dem Fab- oder dem Fab¹-Fragment hiervon enthalten ist.
- 45 **51.** Verfahren nach einem der Ansprüche 47 bis 50, weiterhin umfassend die Phosphorylierung des monoklonalen Antikörpers.
- 50 **52.** Verfahren nach einem der Ansprüche 47 bis 51, weiter umfassend eine radioaktive Markierung des monoklonalen Antikörpers.
- 55 **53.** Verfahren nach Anspruch 1, wobei das Protein aus einem Zytokin besteht.
- 60 **54.** Verfahren nach Anspruch 1, wobei das Protein aus dem Tumornekrose-Faktor (TNF) oder Interleukin-3 (IL-3) besteht.
- 65 **55.** Verfahren nach Anspruch 54, wobei das Protein aus der Gruppe TNF- α und TNF- β ausgewählt ist.
- 70 **56.** Verfahren zur Herstellung eines phosphorylierbaren Proteins aus einem ansonst nicht phosphorylierbaren Protein, umfassend folgende Stufen:

 Identifizieren einer durch eine Proteinkinase (im folgenden als "PK" bezeichnet) erkennbaren Aminosäuresequenz;

50 Definieren und konstruieren einer Nukleotidsequenz mit Kodierung für die Aminosäure-PK-Erkennungssequenz;

 Bestimmen einer Nukleotidsequenz mit Kodierung für das nicht phosphorylierbare Protein;

55 Konstruieren einer modifizierten Nukleotidsequenz mit Kodierung für das phosphorylierbare Protein und

 Expimieren der modifizierten Nukleotidsequenz zur Produktion des phosphorylierbaren Proteins.

57. Verfahren nach Anspruch 56, wobei die Stufe der Konstruktion der modifizierten Nukleotidsequenz ein Anhängen der Nukleotidsequenz mit Kodierung für die Aminosäure-PK-Erkennungssequenz an die Nukleotidsequenz mit Kodierung für das nicht phosphorylierbare Protein beinhaltet.
- 5 58. Verfahren nach Anspruch 56, wobei die Stufe der Konstruktion der modifizierten Nukleotidsequenz eine Einfügung der Nukleotidsequenz mit Kodierung für die Aminosäure-PK-Erkennungssequenz in die Nukleotidsequenz mit Kodierung für das nicht phosphorylierbare Protein beinhaltet.
- 10 59. Verfahren nach Anspruch 56, wobei die in der Identifizierungsstufe identifizierte Aminosäure-PK-Erkennungssequenz Ser enthält.
60. Verfahren nach Anspruch 56, wobei die in der Identifizierungsstufe identifizierte Aminosäure-PK-Erkennungssequenz eine Aminosäure, ausgewählt aus der Gruppe Ser, Thr oder Tyr enthält.
- 15 61. Verfahren nach Anspruch 56, wobei die in der Identifizierungsstufe identifizierte Aminosäure-PK-Erkennungssequenz aus Arg-Arg-Ala-Ser- besteht.
62. Verfahren nach Anspruch 56, wobei die in der Identifizierungsstufe identifizierte Aminosäure-PK-Erkennungssequenz Arg-Arg-Ala-Ser-Xaa, mit Xaa ausgewählt aus der Gruppe sämtlicher Aminosäuren, enthält.
- 20 63. Verfahren nach Anspruch 56, wobei es sich bei der Proteinkinase der Identifizierungsstufe um die cAMP-abhängige Proteinkinase handelt.
- 25 64. Verfahren nach Anspruch 56, wobei das phosphorylierbare Protein mit einer radioaktiven Markierung phosphoryliert ist.

Patentansprüche für folgenden Vertragsstaat : GR

- 30 1. Verfahren zur Herstellung eines Proteins, welches derart modifiziert ist, daß es eine durch eine Proteinkinase erkennbare, als Teil des Proteins phosphorylierbare und in dem nicht modifizierten Protein nicht vorhandene oder für die Phosphorylierung nicht verfügbare Aminosäuresequenz umfaßt.
- 35 2. Verfahren nach Anspruch 1, wobei für die Modifikation durch rekombinante DNA-Techniken gesorgt wurde.
3. Verfahren nach Anspruch 2, wobei es sich bei dem Protein um ein funktionelles Protein handelt.
4. Verfahren nach Anspruch 2, wobei es sich bei dem Protein um ein interferonartiges Protein, ausgewählt aus der Gruppe ein interferonartiges Hu-IFN- α Protein oder ein interferonartiges Hu-IFN- β Protein handelt.
- 40 5. Verfahren nach Anspruch 4, wobei das Protein Hu-IFN- α -artig ist.
6. Verfahren nach Anspruch 4, wobei das Protein Hu-IFN- β -artig ist.
- 45 7. Verfahren nach Anspruch 4, wobei es sich bei dem interferonartigen Hu-IFN- α Protein um ein fusioniertes Hu-IFN- α /gamma Protein handelt.
8. Verfahren nach Anspruch 7, wobei das Hu-IFN- β -artige Protein die schematisch in Figur 3 dargestellte Aminosäuresequenz aufweist.
- 50 9. Verfahren nach Anspruch 7, wobei das Hu-IFN- α -artige Protein aus der Gruppe der in Figur 8 dargestellten und als α -A-P1, α -A-P2 und α -A-P3 identifizierten drei Aminosäuresequenzen ausgewählt ist.
- 55 10. Verfahren nach Anspruch 7, wobei das interferonartige Hu-IFN- α Protein eine biologische Eigenschaft von Hu-IFN- α und/oder von Hu-IFN-gamma aufweist.
11. Verfahren nach Anspruch 4, wobei das Hu-IFN- α -artige Protein eine biologische Eigenschaft von Hu-IFN- α und/oder

von Hu-IFN- β aufweist.

12. Verfahren nach Anspruch 4, wobei das Hu-IFN- α -artige Protein oder Hu-IFN- β -artige Protein eine durch eine Proteinkinase erkennbare Aminosäurekonsenssequenz umfaßt.

13. Verfahren nach Anspruch 12, wobei das Hu-IFN- α -artige Protein eine ein Insert in ein phosphorylierbares Hu-IFN- α -artiges Protein darstellende Aminosäurekonsenssequenz umfaßt.

14. Verfahren nach Anspruch 12, wobei bei dem Hu-IFN- α -artigen Protein die durch die Proteinkinase erkennbare Aminosäurekonsenssequenz mit dem Carboxylende der Aminosäuresequenz von Hu-IFN- α anfusioniert ist.

15. Verfahren nach Anspruch 12, wobei das interferonartige Hu-IFN- α Protein mindestens zwei Aminosäuresegmente, nämlich ein erstes segment, bei dem es sich um die Aminosäuresequenz von natürlichem Hu-IFN- α handelt, und ein zweites Aminosäuresegment, das die durch Proteinkinase erkennbare Aminosäurekonsenssequenz enthält, aufweist, und wobei das zweite Segment in das erste Aminosäuresegment eingebaut oder mit diesem verbunden sein kann.

16. Verfahren nach Anspruch 5, wobei das Hu-IFN- α -artige Protein phosphoryliert ist und ein serumstables Phosphat enthält.

17. Verfahren nach Anspruch 5, wobei das Hu-IFN- α -artige Protein nach der Phosphorylierung in Humanserum stabil ist.

18. Verfahren nach Anspruch 12, wobei die Aminosäurekonsenssequenz des Hu-IFN- α -artigen Proteins Ser enthält.

19. Verfahren nach Anspruch 12, wobei die Aminosäurekonsenssequenz des Hu-IFN- α -artigen Proteins eine Aminosäure, ausgewählt aus der Gruppe Ser, Thr oder Tyr, enthält.

20. Verfahren nach Anspruch 12, wobei die Aminosäurekonsenssequenz des Hu-IFN- α -artigen Proteins aus Arg-Arg-Ala-Ser- besteht.

21. Verfahren nach Anspruch 12, wobei die Aminosäurekonsenssequenz des Hu-IFN- α -artigen Proteins Arg-Arg-Ala-Ser-Xaa mit Xaa ausgewählt aus der Gruppe Gln, Val oder Leu enthält.

22. Verfahren nach Anspruch 12, wobei das Protein aus einem Hu-IFN- α -artigen Protein und die Proteinkinase aus der cAMP-abhängigen Proteinkinase bestehen.

23. Verfahren nach Anspruch 4, weiterhin umfassend die Reinigung des phosphorylierbaren Hu-IFN- α -artigen Proteins.

24. Verfahren nach Anspruch 4, weiter umfassend die Phosphorylierung des phosphorylierbaren Hu-IFN- α -artigen Proteins mit einer radioaktiven Markierung.

25. Verfahren nach Anspruch 24, wobei es sich bei der Markierung um ein β -Strahlung emittierendes Atom handelt.

26. Verfahren nach Anspruch 25, wobei das Atom aus der Gruppe Phosphor und Schwefel ausgewählt ist.

27. Verfahren nach Anspruch 26, wobei es sich bei der radioaktiven Markierung um ^{32}P oder ^{33}P handelt.

28. Verfahren zur Herstellung eines Hu-IFN- α -artigen Proteins oder eines Hu-IFN- β -artigen Proteins, wobei das phosphorylierte Protein der Definition gemäß einem der Ansprüche 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21 oder 23 entspricht und das Verfahren folgende Stufen umfaßt:

(a) Einführen einer durch eine Proteinkinase zur Phosphorylierung erkennbaren Aminosäuresequenz in ein nicht phosphorylierbares Protein oder

(b) Synthese des phosphorylierbaren Proteins aus Aminosäurevorläufern.

29. Verfahren zur Herstellung des phosphorylierbaren Hu-IFN- α -artigen Proteins mit der in Figur 3 dargestellten Ami-

nosäuresequenz oder einer der als α A-P1, α A-P2 und α A-P3 identifizierten und in Figur 8 dargestellten Aminosäuresequenzen, wobei das Verfahren folgende Stufen umfaßt:

(a) Einführen einer durch eine Proteinkinase zur Phosphorylierung erkennbaren Aminosäuresequenz in ein nicht phosphorylierbares Protein oder

(b) Synthese des phosphorylierbaren Proteins aus Aminosäurevorläufern.

30. Verfahren zur Herstellung eines phosphorylierbaren Hu-IFN- β -artigen Proteins des phosphorylierbaren Proteins gemäß Anspruch 6 wobei das verfahren folgende Stufen umfaßt:

(a) Einführen einer durch eine Proteinkinase zur Phosphorylierung erkennbaren Aminosäuresequenz in ein nicht phosphorylierbaren Protein oder

(b) Synthese des phosphorylierbaren Proteins aus Aminosäurevorläufern.

31. Rekombinante DNA-Sequenz mit Kodierung für ein modifiziertes Protein nach Anspruch 1.

32. Rekombinante DNA-Sequenz nach Anspruch 31, enthaltend eine DNA-Sequenz mit Kodierung für ein funktionelles Protein, welches normalerweise nicht phosphorylierbar ist, und eine DNA-Sequenz mit Kodierung für die Aminosäurekonsenssequenz, die durch eine Proteinkinase erkennbar ist und die mutmaßliche Markierungsstelle enthält.

33. Rekombinante DNA-Sequenz nach Anspruch 32, wobei es sich bei dem funktionellen Protein um ein interferonartiges Protein aus der Gruppe Hu-IFN- α oder Hu-IFN- β handelt.

34. Rekombinante DNA-Sequenz nach Anspruch 32, wobei die zweite Nukleotidsequenz ein mit der ersten Nukleotidsequenz gemeinsames Kodon aufweist.

35. Rekombinante DNA-Sequenz, ausgewählt aus der Gruppe DNA-Sequenzen mit Kodierung für ein Protein, ausgewählt aus der Gruppe Hu-IFN- α -artiges Protein oder Hu-IFN- β -artiges Protein, wobei das Protein derart modifiziert wurde, daß es eine durch eine Proteinkinase erkennbare Aminosäurekonsenssequenz mit einer mutmaßlichen phosphorylierbaren Stelle enthält.

36. Rekombinante DNA-Sequenz nach Anspruch 35 mit Kodierung für ein phosphorylierbares Hu-IFN- α -artiges Protein.

37. Rekombinanter DNA-Klonierungsvektor mit der Eignung zum Exprimieren eines exogenen funktionellen Proteins, wobei der Vektor die DNA-Sequenz mit Kodierung für ein Hu-IFN- α -artiges Protein oder ein Hu-IFN- β -artiges Protein beinhaltet, das Protein derart modifiziert wurde, daß es eine durch eine Proteinkinase erkennbare Aminosäurekonsenssequenz mit einer mutmaßlichen Phosphorylierungsstelle enthält und das phosphorylierbare Protein aus einem interferonartigen Hu-IFN- α Protein oder einem interferonartigen Hu-IFN- β Protein besteht.

38. Klonierungsvektor nach Anspruch 37, wobei es sich bei dem kodierten Protein um das interferonartige Protein Hu-IFN- α handelt.

39. Klonierungsvektor nach Anspruch 37, wobei es sich bei dem kodierten Protein um das interferonartige Protein Hu-IFN- β handelt.

40. Klonierungsvektor nach Anspruch 37, enthaltend die DNA-Sequenz mit Kodierung für ein fusioniertes Hu-IFN- α /gamma Protein.

41. Klonierungsvektor nach Anspruch 37, enthaltend die DNA-Sequenz mit Kodierung für die Figur 8 mit -P1, -P2 und bzw. -P3 bezeichneten modifizierten Interferone.

42. Transformierte Wirtszelle mit der Fähigkeit zum Exprimieren des funktionellen Proteins, wobei die Wirtszelle einen rekombinanten DNA-Vektor mit Kodierung für ein Hu-IFN- α -artiges Protein oder ein Hu-IFN- β -artiges Protein, welches derart modifiziert wurde, daß es eine durch eine Proteinkinase erkennbare Aminosäurekonsenssequenz enthält, enthält.

- 5
43. Verfahren zur Zubereitung einer biologisch aktiven Zusammensetzung, umfassend die Stufe des Vermischen eines biologisch verträglichen Trägers und einer biologisch wirksamen Menge eines phosphorylierten Proteins entsprechend der Definition gemäß einem der Ansprüche 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26 oder 27.
- 10
44. Verfahren nach Anspruch 1, wobei das phosphorylierbare Protein eine biologische Eigenschaft des nicht phosphorylierbaren Proteins aufweist.
- 15
45. Verfahren nach Anspruch 1, wobei das phosphorylierbare Protein mindestens zwei Aminosäuresegmente, nämlich ein erstes Segment, bei dem es sich um die Aminosäuresequenz des nicht phosphorylierbaren Proteins handelt, und ein zweites Aminosäuresegment, welches die durch die Proteinkinase erkennbare Aminosäuresequenz enthält, wobei das zweite Segment in das erste Aminosäuresegment eingebaut oder mit diesem verbunden sein kann.
- 20
46. Verfahren nach Anspruch 1, wobei das phosphorylierbare Protein phosphoryliert ist und ein serumstabiles Phosphat enthält.
- 25
47. Verfahren nach Anspruch 1, wobei das Protein aus einem phosphorylierbaren monoklonalen Antikörper besteht.
- 30
48. Verfahren nach Anspruch 47, wobei der monoklonale Antikörper aus einer Chimäre, einem Hybrid oder modifizierten monoklonalen Antikörper besteht.
- 35
49. Verfahren nach Anspruch 47, wobei der monoklonale Antikörper Tumor-spezifisch ist.
- 40
50. Verfahren nach Anspruch 47, wobei die Phosphorylierungsstelle in dem Fab- oder dem Fab¹-Fragment hiervon enthalten ist.
- 45
51. Verfahren nach einem der Ansprüche 47 bis 50, weiterhin umfassend die Phosphorylierung des monoklonalen Antikörpers.
- 50
52. Verfahren nach einem der Ansprüche 47 bis 51, weiter umfassend eine radioaktive Markierung des monoklonalen Antikörpers.
- 55
53. Verfahren nach Anspruch 1, wobei das Protein aus einem Zytokin besteht.
56. Verfahren nach Anspruch 1, wobei das Protein aus dem Tumornekrose-Faktor (TNF) oder Interleukin-3 (IL-3) besteht.
57. Verfahren nach Anspruch 54, wobei das Protein aus der Gruppe TNF- α und TNF- β ausgewählt ist.
58. Verfahren zur Herstellung eines phosphorylierbaren Proteins aus einem ansonst nicht phosphorylierbaren Protein, umfassend folgende Stufen:
- Identifizieren einer durch eine Proteinkinase (im folgenden als "PK" bezeichnet) erkennbaren Aminosäuresequenz;
- Definieren und konstruieren einer Nukleotidsequenz mit Kodierung für die Aminosäure-PK-Erkennungssequenz;
- Bestimmen einer Nukleotidsequenz mit Kodierung für das nicht phosphorylierbare Protein;
- Konstruieren einer modifizierten Nukleotidsequenz mit Kodierung für das phosphorylierbare Protein und
- Exprimieren der modifizierten Nukleotidsequenz zur Produktion des phosphorylierbaren Proteins.
59. Verfahren nach Anspruch 58, wobei die Stufe der Konstruktion der modifizierten Nukleotidsequenz ein Anhängen der Nukleotidsequenz mit Kodierung für die Aminosäure-PK-Erkennungssequenz an die Nukleotidsequenz mit Kodierung für das nicht phosphorylierbare Protein beinhaltet.

58. Verfahren nach Anspruch 56, wobei die Stufe der Konstruktion der modifizierten Nukleotidsequenz eine Einfügung der Nukleotidsequenz mit Kodierung für die Aminosäure-PK-Erkennungssequenz in die Nukleotidsequenz mit Kodierung für das nicht phosphorylierbare Protein beinhaltet.
- 5 59. Verfahren nach Anspruch 56, wobei die in der Identifizierungsstufe identifizierte Aminosäure-PK-Erkennungssequenz Ser enthält.
60. Verfahren nach Anspruch 56, wobei die in der Identifizierungsstufe identifizierte Aminosäure-PK-Erkennungssequenz eine Aminosäure, ausgewählt aus der Gruppe Ser, Thr oder Tyr enthält.
- 10 61. Verfahren nach Anspruch 56, wobei die in der Identifizierungsstufe identifizierte Aminosäure-PK-Erkennungssequenz aus Arg-Arg-Ala-Ser- besteht.
62. Verfahren nach Anspruch 56, wobei die in der Identifizierungsstufe identifizierte Aminosäure-PK-Erkennungssequenz Arg-Arg-Ala-Ser-Xaa, mit Xaa ausgewählt aus der Gruppe sämtlicher Aminosäuren, enthält.
- 15 63. Verfahren nach Anspruch 56, wobei es sich bei der Proteinkinase der Identifizierungsstufe um die cAMP-abhängige Proteinkinase handelt.
- 20 64. Verfahren nach Anspruch 56, wobei das phosphorylierbare Protein mit einer radioaktiven Markierung phosphoryliert ist.

Revendications

Revendications pour les Etats contractants suivants : DE, FR, GB, IT, NL, SE, CH, BE, LU, AT

- 30 1. Protéine qui est modifiée pour comprendre une séquence d'acides aminés reconnaissable par une protéinekinase, laquelle séquence d'acides aminés est, en tant que partie de la protéine, phosphorylable et n'était pas présente, ou n'était pas disponible pour la phosphorylation, dans la protéine non modifiée.
2. Protéine phosphorylable selon la revendication 1, dans laquelle ladite modification est fournie par des techniques d'ADN recombinant.
- 35 3. Protéine phosphorylable selon la revendication 2, qui est une protéine fonctionnelle.
4. Protéine phosphorylable selon la revendication 2, qui est une protéine semblable à un interféron choisie dans le groupe constitué d'une protéine Hu-IFN- α semblable à un interféron et d'une protéine Hu-IFN- β semblable à un interféron.
- 40 5. Protéine phosphorylable semblable à Hu-IFN- α selon la revendication 4.
6. Protéine phosphorylable semblable Hu-IFN- β selon la revendication 4.
- 45 7. Hu-IFN- α phosphorylable selon la revendication 4, dans laquelle la protéine Hu-IFN- α semblable à un interféron est une protéine fusionnée Hu-IFN- α /gamma.
8. Protéine phosphorylable semblable à Hu-IFN- β fusionnée selon la revendication 7, qui a la séquence d'acides aminés présentée schématiquement à la figure 3.
- 50 9. Protéine phosphorylable semblable à Hu-IFN- α selon la revendication 5, qui est choisie dans le groupe des trois séquences d'acides aminés présentées à la figure 8 et identifiées comme α A-P1, α A-P2 et α A-P3.
- 55 10. Hu-IFN- α phosphorylable selon la revendication 7, dans lequel la protéine Hu-IFN- α semblable à un interféron possède une propriété biologique de Hu-IFN- α et/ou de Hu-IFN-gamma.
11. Protéine phosphorylable semblable à Hu-IFN- α selon la revendication 4, qui possède une propriété biologique de

Hu-IFN- α et/ou de Hu-IFN- β .

12. Protéine phosphorylable semblable à Hu-IFN- α ou semblable à Hu-IFN- β selon la revendication 4, qui comprend une séquence consensus d'acides aminés qui est reconnaissable par une protéinekinase.
13. Protéine phosphorylable semblable à Hu-IFN- α selon la revendication 12, qui comprend une séquence consensus d'acides aminés qui est un segment d'insertion dans une protéine phosphorylable semblable à Hu-IFN- α ,
14. Protéine phosphorylable semblable à Hu-IFN- α selon la revendication 12, dans laquelle la séquence consensus d'acides aminés reconnaissable par la protéinekinase est fusionnée à l'extrémité carboxyle de la séquence d'acides aminés de Hu-IFN- α .
15. Protéine phosphorylable Hu-IFN- α semblable à un interféron selon la revendication 12, qui contient au moins deux segments d'acides aminés, un premier segment qui est la séquence d'acides aminés de Hu-IFN- α naturel et un second segment d'acides aminés qui contient la séquence consensus d'acides aminés qui est reconnaissable par une protéinekinase, ce second segment pouvant être incorporé dans le premier segment d'acides aminés ou lié à celui-ci.
16. Protéine phosphorylable semblable à Hu-IFN- α selon la revendication 5, qui est phosphorylée et contient un phosphate stable dans un sérum.
17. Protéine phosphorylable semblable à Hu-IFN- α selon la revendication 5, qui, après phosphorylation, est stable dans du sérum humain.
18. Protéine phosphorylable semblable à Hu-IFN- α selon la revendication 12, dans laquelle la séquence consensus d'acides aminés contient Ser.
19. Protéine phosphorylable semblable à Hu-IFN- α selon la revendication 12, dans laquelle la séquence consensus d'acides aminés contient un acide aminé choisi dans le groupe constitué de Ser, Thr ou Tyr.
20. Protéine phosphorylable semblable à Hu-IFN- α selon la revendication 12, dans laquelle la séquence consensus d'acides aminés est Arg-Arg-Ala-Ser-.
21. Protéine phosphorylable semblable à Hu-IFN- α selon la revendication 12, dans laquelle la séquence consensus d'acides aminés contient Arg-Arg-Ala-Ser-Xaa, où Xaa est choisi dans le groupe de Gln, Val ou Leu.
22. Protéine phosphorylable semblable à Hu-IFN- α selon la revendication 12, dans laquelle la protéinekinase est la protéinekinase dépendante de l'AMPc.
23. Protéine phosphorylable semblable à Hu-IFN- α selon la revendication 4, qui est purifiée.
24. Protéine phosphorylable semblable à Hu-IFN- α selon la revendication 4, qui est phosphorylée avec un marqueur radioactif.
25. Protéine phosphorylable semblable à Hu-IFN- α selon la revendication 24, dans laquelle le marqueur est un atome émettant des rayons bêta.
26. Protéine phosphorylable semblable à Hu-IFN- α selon la revendication 25, dans laquelle l'atome est choisi dans le groupe constitué du phosphore et du soufre.
27. Protéine phosphorylable semblable à Hu-IFN- α selon la revendication 26, dans laquelle le marqueur radioactif est ^{32}P ou ^{33}P .
28. Protéine semblable à Hu-IFN- α ou protéine semblable à Hu-IFN- β phosphorylable, la protéine phosphorylée étant définie dans l'une quelconque des revendications 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21 ou 23.
29. Protéine phosphorylable semblable à Hu-IFN- α , ayant la séquence d'acides aminés présentée à la figure 3 ou l'une des séquences d'acides aminés identifiées comme $\alpha\text{A-p1}$, $\alpha\text{A-P2}$ et $\alpha\text{A-P3}$, présentées à la figure 8.

30. Protéine phosphorylable semblable à Hu-IFN- β de la protéine phosphorylée définie dans la revendication 6.
31. Séquence d'ADN recombinant qui code pour une protéine modifiée selon la revendication 1.
- 5 32. Séquence d'ADN recombinant selon la revendication 31, qui contient une séquence d'ADN qui code pour une protéine fonctionnelle qui normalement n'est pas phosphorylable et une séquence d'ADN qui code pour la séquence consensus d'acides aminés qui est reconnaissable par une protéinekinase et qui contient le site de marquage supposé.
- 10 33. Séquence d'ADN recombinant selon la revendication 32, dans laquelle la protéine fonctionnelle est une protéine semblable à un interféron choisie dans le groupe constitué de Hu-IFN- α ou de Hu-IFN- β .
34. Séquence d'ADN recombinant selon la revendication 32, dans laquelle la seconde séquence de nucléotides a un codon qui est commun avec la première séquence de nucléotides.
- 15 35. Séquence d'ADN recombinant, qui est choisie dans le groupe constitué des séquences d'ADN codant pour une protéine choisie dans le groupe constitué d'une protéine semblable à Hu-IFN- α ou d'une protéine semblable à Hu-IFN- β , cette protéine ayant été modifiée pour contenir une séquence consensus d'acides aminés reconnaissable par une protéinekinase, comprenant un site phosphorylable supposé.
- 20 36. Séquence d'ADN - recombinant selon la revendication 35, qui code pour une protéine phosphorylable semblable à Hu-IFN- α .
- 25 37. Vecteur de clonage d'ADN recombinant utile pour exprimer une protéine fonctionnelle exogène, ce vecteur incluant la séquence d'ADN qui code pour une protéine semblable à Hu-IFN- α ou une protéine semblable à Hu-IFN- β , cette protéine ayant été modifiée pour contenir une séquence consensus d'acides aminés reconnaissable par une protéinekinase, comprenant un site de phosphorylation supposé, la protéine phosphorylable étant une protéine Hu-IFN- α semblable à un interféron ou une protéine Hu-IFN- β semblable à un interféron.
- 30 38. Vecteur de clonage selon la revendication 37, dans lequel la protéine codée est la protéine Hu-IFN- α semblable à un interféron.
39. Vecteur de clonage selon la revendication 37, dans lequel la protéine codée est la protéine Hu-IFN- β semblable à un interféron.
- 35 40. Vecteur de clonage selon la revendication 37, qui contient la séquence d'ADN codant pour une protéine fusionnée Hu-IFN- α /gamma.
41. Vecteur de clonage selon la revendication 37, qui contient la séquence d'ADN codant pour les interférons modifiés dénommés -P1, -P2 et -P3 dans la figure 8, respectivement.
- 40 42. Cellule hôte transformée capable d'exprimer une protéine fonctionnelle, cette cellule hôte contenant un vecteur d'ADN recombinant codant pour une protéine semblable à Hu-IFN- α ou une protéine semblable à Hu-IFN- β qui a été modifiée pour contenir une séquence consensus d'acides aminés reconnaissable par une protéinekinase.
- 45 43. Composition biologiquement active, qui comprend un véhicule biologiquement compatible et, en une quantité biologique efficace, une protéine phosphorylée définie dans l'une quelconque des revendications 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26 ou 27.
- 50 44. Protéine phosphorylable selon la revendication 1, qui possède une propriété biologique de la protéine non phosphorylable.
- 45 45. Protéine phosphorylable selon la revendication 1, qui contient au moins deux segments d'acides aminés, un premier segment qui est la séquence d'acides aminés de la protéine non phosphorylable et un second segment d'acides aminés qui contient la séquence d'acides aminés reconnaissable par la protéinekinase, ce second segment pouvant être incorporé dans le premier segment d'acides aminés ou lié à celui-ci.
- 55 46. Protéine phosphorylable selon la revendication 1, qui est phosphorylée et contient un phosphate stable dans un

sérum.

47. Protéine selon la revendication 1, qui est un anticorps monoclonal phosphorylable.

48. Anticorps monoclonal selon la revendication 47, qui est un anticorps monoclonal chimérique, hybride ou modifié.

49. Anticorps monoclonal selon la revendication 47, qui est spécifique à une tumeur.

50. Anticorps monoclonal selon la revendication 47, dans lequel le site pour la phosphorylation est dans le fragment Fab ou Fab¹ de celui-ci.

51. Anticorps monoclonal selon l'une quelconque des revendications 47 à 50, qui est phosphorylé.

52. Anticorps monoclonal selon l'une quelconque des revendications 47 à 51, qui est radiomarké.

53. Protéine selon la revendication 1, qui est une cytokine.

54. Protéine selon la revendication 1, qui est le facteur de nécrose des tumeurs (TNF) ou l'interleukine-3 (IL-3).

55. Protéine selon la revendication 54, qui est choisie dans le groupe constitué du TNF- α et du TNF- β .

56. Procédé pour préparer une protéine phosphorylable à partir d'une protéine par ailleurs non phosphorylable, comprenant les étapes de :

identification d'une séquence d'acides aminés reconnaissable par une protéinekinase (ci-après "PK") ;
définition et construction d'une séquence de nucléotides qui code pour ladite séquence d'acides aminés reconnaissable par une PK ;
détermination d'une séquence de nucléotides codant pour ladite protéine non phosphorylable ;
construction d'une séquence de nucléotides modifiée codant pour ladite protéine phosphorylable ;
expression de ladite séquence de nucléotides modifiée pour produire ladite protéine phosphorylable.

57. Procédé selon la revendication 56, dans lequel ladite étape de construction de ladite séquence de nucléotides modifiée inclut la fixation de ladite séquence de nucléotides codant pour ladite séquence d'acides aminés reconnaissable par une PK à ladite séquence de nucléotides codant pour ladite protéine non phosphorylable.

58. Procédé selon la revendication 56, dans lequel ladite étape de construction de ladite séquence de nucléotides modifiée inclut l'introduction de ladite séquence de nucléotides codant pour ladite séquence d'acides aminés reconnaissable par une PK dans ladite séquence de nucléotides codant pour ladite protéine non phosphorylable.

59. Procédé selon la revendication 56, dans lequel ladite séquence d'acides aminés reconnaissable par une PK, identifiée dans ladite étape d'identification, contient Ser.

60. Procédé selon la revendication 56, dans lequel ladite séquence d'acides aminés reconnaissable par une PK, identifiée dans ladite étape d'identification, contient un acide aminé choisi dans le groupe constitué de Ser, Thr ou Tyr.

61. Procédé selon la revendication 56, dans lequel ladite séquence d'acides aminés reconnaissable par une PK identifiée dans ladite étape d'identification est Arg-Arg-Ala-Ser-.

62. Procédé selon la revendication 56, dans lequel ladite séquence d'acides aminés reconnaissable par une PK identifiée dans ladite étape d'identification contient Arg-Arg-Ala-Ser-Xaa, où Xaa est choisi dans le groupe de tous les acides aminés.

63. Procédé selon la revendication 56, dans lequel la protéinekinase de ladite étape d'identification est la protéinekinase dépendant de l'AMPc.

64. Procédé selon la revendication 56, dans lequel ladite protéine phosphorylable est phosphorylée avec un marqueur radioactif.

Revendications pour l'Etat contractant suivant : ES

- 5 1. Procédé de préparation d'une protéine qui est modifiée pour comprendre une séquence d'acides aminés reconnaissable par une protéinekinase, laquelle séquence d'acides aminés est, en tant que partie de la protéine, phosphorylable et n'était pas présente, ou n'était pas disponible pour la phosphorylation, dans la protéine non modifiée.
2. Procédé selon la revendication 1, dans laquelle ladite modification est fournie par des techniques d'ADN recombinant.
- 10 3. Procédé selon la revendication 2, qui est une protéine fonctionnelle.
4. Procédé selon la revendication 2, qui est une protéine semblable à un interféron choisie dans le groupe constitué d'une- protéine Hu-IFN- α semblable à un interféron et d'une protéine Hu-IFN- β semblable à un interféron.
- 15 5. Procédé selon la revendication 4, dans lequel la protéine est semblable à Hu-IFN- α .
6. Procédé selon la revendication 4, dans lequel la protéine est semblable à Hu-IFN- β .
- 20 7. Procédé selon la revendication 4, dans lequel la protéine Hu-IFN- α semblable à un interféron est une protéine fusionnée Hu-IFN- α /gamma.
8. Procédé selon la revendication 7, dans lequel la protéine semblable à Hu-IFN- β a la séquence d'acides aminés présentée schématiquement à la figure 3.
- 25 9. Procédé selon la revendication 7, dans lequel la protéine semblable à Hu-IFN- α est choisie dans le groupe des trois séquences d'acides aminés présentées à la figure 8 et identifiées comme α A-P1, α A-P2 et α A-P3.
- 30 10. Procédé selon la revendication 7, dans lequel la protéine Hu-IFN- α semblable à un interféron possède une propriété biologique de Hu-IFN- α et/ou de Hu-IFN-gamma.
11. Procédé selon la revendication 4, dans lequel la protéine semblable à Hu-IFN- α possède une propriété biologique de Hu-IFN- α et/ou de Hu-IFN- β .
- 35 12. Procédé selon la revendication 4, dans lequel la protéine semblable à Hu-IFN- α ou semblable à Hu-IFN- β comprend une séquence consensus d'acides aminés qui est reconnaissable par une protéinekinase.
13. Procédé selon la revendication 12, dans lequel la protéine semblable à Hu-IFN- α comprend une séquence consensus d'acides aminés qui est un segment d'insertion dans une protéine phosphorylable semblable à Hu-IFN- α .
- 40 14. Procédé selon la revendication 12, dans lequel la protéine semblable à Hu-IFN- α comprenant la séquence consensus d'acides aminés reconnaissable par la protéinekinase est fusionnée à l'extrémité carboxyle de la séquence d'acides aminés de Hu-IFN- α .
- 45 15. Procédé selon la revendication 12, dans lequel la protéine Hu-IFN- α semblable à un interféron contient au moins deux segments d'acides aminés, un premier segment qui est la séquence d'acides aminés de Hu-IFN- α naturel et un second segment d'acides aminés qui contient la séquence consensus d'acides aminés qui est reconnaissable par une protéinekinase, ce second segment pouvant être incorporé dans le premier segment d'acides aminés ou lié à celui-ci.
- 50 16. Procédé selon la revendication 5, dans lequel la protéine semblable à Hu-IFN- α est phosphorylée et contient un phosphate stable dans un sérum.
17. Procédé selon la revendication 5, dans lequel la protéine semblable à Hu-IFN- α , après phosphorylation, est stable dans du sérum humain.
- 55 18. Procédé selon la revendication 12, dans lequel la séquence consensus d'acides aminés de la protéine semblable à Hu-IFN- α contient Ser.

19. Procédé selon la revendication 12, dans lequel la séquence consensus d'acides aminés de la protéine semblable à Hu-IFN- α contient un acide aminé choisi dans le groupe constitué de Ser, Thr ou Tyr.
- 5 20. Procédé selon la revendication 12, dans lequel la séquence consensus d'acides aminés de la protéine semblable à Hu-IFN- α est Arg-Arg-Ala-Ser-.
21. Procédé selon la revendication 12, dans lequel la séquence consensus d'acides aminés de la protéine semblable à Hu-IFN- α contient Arg-Arg-Ala-Ser-Xaa, où Xaa est choisi dans le groupe de Gln, Val ou Leu.
- 10 22. Procédé selon la revendication 12, dans lequel la protéine est la protéine semblable à Hu-IFN- α et la protéine kinase est la protéine kinase dépendante de l'AMPc.
23. Procédé selon la revendication 4, comprenant en outre la purification de la protéine phosphorylable semblable à Hu-IFN- α .
- 15 24. Procédé selon la revendication 4, comprenant en outre la phosphorylation de la protéine semblable à Hu-IFN- α avec un marqueur radioactif.
25. Procédé selon la revendication 24, dans lequel le marqueur est un atome émettant des rayons bêta.
- 20 26. Procédé selon la revendication 25, dans lequel l'atome est choisi dans le groupe constitué du phosphore et du soufre.
27. Procédé selon la revendication 26, dans lequel le marqueur radioactif est ^{32}P ou ^{33}P .
- 25 28. Procédé de préparation d'une protéine semblable à Hu-IFN- α ou d'une protéine semblable à Hu-IFN- β , la protéine phosphorylée étant définie dans l'une quelconque des revendications 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21 ou 23, le procédé comprenant l'étape de (a) introduction, dans une protéine qui n'est pas phosphorylable, d'une séquence d'acides aminés reconnaissable par une protéine kinase pour la phosphorylation ou de (b) synthèse de ladite protéine phosphorylable à partir de précurseurs d'acides aminés.
- 30 29. Procédé de préparation de la protéine phosphorylable semblable à Hu-IFN- α ayant la séquence d'acides aminés présentée à la figure 3 ou l'une des séquences d'acides aminés identifiées comme $\alpha\text{A-P1}$, $\alpha\text{A-P2}$ et $\alpha\text{A-P3}$, présentées à la figure 8, le procédé comprenant l'étape de (a) introduction, dans une protéine qui n'est pas phosphorylable, d'une séquence d'acides aminés reconnaissable par une protéine kinase pour la phosphorylation ou de (b) synthèse de ladite protéine phosphorylable à partir de précurseurs d'acides aminés.
- 35 30. Procédé de préparation de la protéine phosphorylable semblable à Hu-IFN- β de la protéine phosphorylée définie dans la revendication 6, le procédé comprenant l'étape de (a) introduction, dans une protéine qui n'est pas phosphorylable, d'une séquence d'acides aminés reconnaissable par une protéine kinase pour la phosphorylation ou de (b) synthèse de ladite protéine phosphorylable à partir de précurseurs d'acides aminés.
- 40 31. Procédé de préparation d'une séquence d'ADN recombinant qui code pour une protéine modifiée telle que définie dans la revendication 1 par des techniques d'ADN recombinant.
- 45 32. Procédé selon la revendication 31, dans lequel la séquence d'ADN recombinant contient une séquence d'ADN qui code pour une protéine fonctionnelle qui normalement n'est pas phosphorylable et une séquence d'ADN qui code pour la séquence consensus d'acides aminés qui est reconnaissable par une protéine kinase et qui contient le site de marquage supposé.
- 50 33. Procédé selon la revendication 32, dans lequel la protéine fonctionnelle est une protéine semblable à un interféron choisie dans le groupe constitué de Hu-IFN- α ou de Hu-IFN- β .
34. Procédé selon la revendication 32, dans lequel la seconde séquence de nucléotides a un codon qui est commun avec la première séquence de nucléotides.
- 55 35. Procédé de préparation d'une séquence d'ADN recombinant qui est choisie dans le groupe constitué des séquences d'ADN codant pour une protéine choisie dans le groupe constitué d'une protéine semblable à Hu-IFN- α ou d'une protéine semblable à Hu-IFN- β , cette protéine ayant été modifiée pour contenir une séquence consensus d'acides

aminés reconnaissable par une protéinekinase, comprenant un site phosphorylable supposé, par des techniques d'ADN recombinant.

- 5 36. Procédé selon la revendication 35, dans lequel la séquence d'ADN recombinant code pour une protéine phosphorylable semblable à Hu-IFN- α .
- 10 37. Procédé de préparation d'un vecteur de clonage d'ADN recombinant utile pour exprimer une protéine fonctionnelle exogène, ce vecteur incluant la séquence d'ADN qui code pour une protéine semblable à Hu-IFN- α ou une protéine semblable à Hu-IFN- β , cette protéine ayant été modifiée pour contenir une séquence consensus d'acides aminés reconnaissable par une protéinekinase, comprenant un site de phosphorylation supposé, la protéine phosphorylable étant une protéine Hu-IFN- α semblable à un interféron ou une protéine Hu-IFN- β semblable à un interféron, par des techniques d'ADN recombinant.
- 15 38. Procédé selon la revendication 37, dans lequel la protéine codée est la protéine Hu-IFN- α semblable à un interféron.
39. Procédé selon la revendication 37, dans lequel la protéine codée est la protéine Hu-IFN- β semblable à un interféron.
- 20 40. Procédé selon la revendication 37, dans lequel le vecteur de clonage contient la séquence d'ADN codant pour une protéine fusionnée Hu-IFN- α /gamma.
41. Procédé selon la revendication 37, dans lequel le vecteur de clonage contient la séquence d'ADN codant pour les interférons modifiés dénommés -P1, -P2 et -P3 dans la figure 8, respectivement.
- 25 42. Procédé de préparation d'une cellule hôte transformée capable d'exprimer une protéine fonctionnelle, cette cellule hôte contenant un vecteur d'ADN recombinant codant pour une protéine semblable à Hu-IFN- α ou une protéine semblable à Hu-IFN- β qui a été modifiée pour contenir une séquence consensus d'acides aminés reconnaissable par une protéinekinase, comprenant la transformation de la cellule hôte avec le vecteur d'ADN recombinant.
- 30 43. Procédé de fabrication d'une composition biologiquement active, comprenant les étapes de mélange d'un véhicule biologiquement compatible et, d'une quantité biologique efficace, d'une protéine phosphorylée telle que définie dans l'une quelconque des revendications 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26 ou 27.
- 35 44. Procédé selon la revendication 1, dans lequel la protéine phosphorylable possède une propriété biologique de la protéine non phosphorylable.
- 40 45. Procédé selon la revendication 1, dans lequel la protéine phosphorylable contient au moins deux segments d'acides aminés, un premier segment qui est la séquence d'acides aminés de la protéine non phosphorylable et un second segment d'acides aminés qui contient la séquence d'acides aminés reconnaissable par la protéinekinase, ce second segment pouvant être incorporé dans le premier segment d'acides aminés ou lié à celui-ci.
46. Procédé selon la revendication 1, dans lequel la protéine phosphorylable est phosphorylée et contient un phosphate stable dans un sérum.
- 45 47. Procédé selon la revendication 1, dans lequel la protéine est un anticorps monoclonal phosphorylable.
48. Procédé selon la revendication 47, dans lequel l'anticorps monoclonal est un anticorps monoclonal chimérique, hybride ou modifié.
- 50 49. Procédé selon la revendication 47, dans lequel l'anticorps monoclonal est spécifique à une tumeur.
50. Procédé selon la revendication 47, dans lequel le site pour la phosphorylation est dans le fragment Fab ou Fab¹ de celui-ci.
- 55 51. Procédé selon l'une quelconque des revendications 47 à 50, comprenant en outre la phosphorylation de l'anticorps monoclonal.
52. Procédé selon l'une quelconque des revendications 47 à 51, comprenant en outre le radiomarquage de l'anticorps

monoclonale

53. Procédé selon la revendication 1, dans lequel la protéine est une cytokine.

5 54. Procédé selon la revendication 1, dans lequel la protéine est le facteur de nécrose des tumeurs (TNF) ou l'interleukine-3 (IL-3).

55. Procédé selon la revendication 54, dans lequel la protéine est choisie dans le groupe constitué du TNF- α et du TNF- β .

10 56. Procédé pour préparer une protéine phosphorylable à partir d'une protéine par ailleurs non phosphorylable, comprenant les étapes de :

15 identification d'une séquence d'acides aminés reconnaissable par une protéine kinase (ci-après "PK") ;
définition et construction d'une séquence de nucléotides qui code pour ladite séquence d'acides aminés reconnaissable par une PK ;
détermination d'une séquence de nucléotides codant pour ladite protéine non phosphorylable ;
construction d'une séquence de nucléotides modifiée codant pour ladite protéine phosphorylable ;
20 expression de ladite séquence de nucléotides modifiée pour produire ladite protéine phosphorylable.

25 57. Procédé selon la revendication 56, dans lequel ladite étape de construction de ladite séquence de nucléotides modifiée inclut la fixation de ladite séquence de nucléotides codant pour ladite séquence d'acides aminés reconnaissable par une PK à ladite séquence de nucléotides codant pour ladite protéine non phosphorylable.

30 58. Procédé selon la revendication 56, dans lequel ladite étape de construction de ladite séquence de nucléotides modifiée inclut l'introduction de ladite séquence de nucléotides codant pour ladite séquence d'acides aminés reconnaissable par une PK dans ladite séquence de nucléotides codant pour ladite protéine non phosphorylable.

35 59. Procédé selon la revendication 56, dans lequel ladite séquence d'acides aminés reconnaissable par une PK, identifiée dans ladite étape d'identification, contient Ser.

60. Procédé selon la revendication 56, dans lequel ladite séquence d'acides aminés reconnaissable par une PK, identifiée dans ladite étape d'identification, contient un acide aminé choisi dans le groupe constitué de Ser, Thr ou Tyr.

40 61. Procédé selon la revendication 56, dans lequel ladite séquence d'acides aminés reconnaissable par une PK identifiée dans ladite étape d'identification est Arg-Arg-Ala-Ser-.

62. Procédé selon la revendication 56, dans lequel ladite séquence d'acides aminés reconnaissable par une PK identifiée dans ladite étape d'identification contient Arg-Arg-Ala-Ser-Xaa, où Xaa est choisi dans le groupe de tous les acides aminés.

63. Procédé selon la revendication 56, dans lequel la protéine kinase de ladite étape d'identification est la protéine kinase dépendant de l'AMPc.

45 64. Procédé selon la revendication 56, dans lequel ladite protéine phosphorylable est phosphorylée avec un marqueur radioactif.

Revendications pour l'Etat contractant suivant : GR

50 1. Procédé de préparation d'une protéine qui est modifiée pour comprendre une séquence d'acides aminés reconnaissable par une protéine kinase, laquelle séquence d'acides aminés est, en tant que partie de la protéine, phosphorylable et n'était pas présente, ou n'était pas disponible pour la phosphorylation, dans la protéine non modifiée

55 2. Procédé selon la revendication 1, dans laquelle ladite modification est fournie par des techniques d'ADN recombinant.

3. Procédé selon la revendication 2, qui est une protéine fonctionnelle.

4. Procédé selon la revendication 2, qui est une protéine semblable à un interféron choisie dans le groupe constitué d'une- protéine Hu-IFN- α semblable à un interféron et d'une protéine Hu-IFN- β semblable à un interféron.
5. Procédé selon la revendication 4, dans lequel la protéine est semblable à Hu-IFN- α .
6. Procédé selon la revendication 4, dans lequel la protéine est semblable à Hu-IFN- β .
7. Procédé selon la revendication 4, dans lequel la protéine Hu-IFN- α semblable à un interféron est une protéine fusionnée Hu-IFN- α /gamma.
8. Procédé selon la revendication 7, dans lequel la protéine semblable à Hu-IFN- β a la séquence d'acides aminés présentée schématiquement à la figure 3.
9. Procédé selon la revendication 7, dans lequel la protéine semblable à Hu-IFN- α est choisie dans le groupe des trois séquences d'acides aminés présentées à la figure 8 et identifiées comme α A-P1, α A-P2 et α A-P3.
10. Procédé selon la revendication 7, dans lequel la protéine Hu-IFN- α semblable à un interféron possède une propriété biologique de Hu-IFN- α et/ou de Hu-IFN-gamma.
11. Procédé selon la revendication 4, dans lequel la protéine semblable à Hu-IFN- α possède une propriété biologique de Hu-IFN- α et/ou de Hu-IFN- β .
12. Procédé selon la revendication 4, dans lequel la protéine semblable à Hu-IFN- α ou semblable à Hu-IFN- β comprend une séquence consensus d'acides aminés qui est reconnaissable par une protéinekinase.
13. Procédé selon la revendication 12, dans lequel la protéine semblable à Hu-IFN- α comprend une séquence consensus d'acides aminés qui est un segment d'insertion dans une protéine phosphorylable semblable à Hu-IFN- α .
14. Procédé selon la revendication 12, dans lequel la protéine semblable à Hu-IFN- α comprenant la séquence consensus d'acides aminés reconnaissable par la protéinekinase est fusionnée à l'extrémité carboxyle de la séquence d'acides aminés de Hu-IFN- α .
15. Procédé selon la revendication 12, dans lequel la protéine Hu-IFN- α semblable à un interféron contient au moins deux segments d'acides aminés, un premier segment qui est la séquence d'acides aminés de Hu-IFN- α naturel et un second segment d'acides aminés qui contient la séquence consensus d'acides aminés qui est reconnaissable par une protéinekinase, ce second segment pouvant être incorporé dans le premier segment d'acides aminés ou lié à celui-ci.
16. Procédé selon la revendication 5, dans lequel la protéine semblable à Hu-IFN- α est phosphorylée et contient un phosphate stable dans un sérum.
17. Procédé selon la revendication 5, dans lequel la protéine semblable à Hu-IFN- α , après phosphorylation, est stable dans du sérum humain.
18. Procédé selon la revendication 12, dans lequel la séquence consensus d'acides aminés de la protéine semblable à Hu-IFN- α contient Ser.
19. Procédé selon la revendication 12, dans lequel la séquence consensus d'acides aminés de la protéine semblable à Hu-IFN- α contient un acide aminé choisi dans le groupe constitué de Ser, Thr ou Tyr.
20. Procédé selon la revendication 12, dans lequel la séquence consensus d'acides aminés de la protéine semblable à Hu-IFN- α est Arg-Arg-Ala-Ser-.
21. Procédé selon la revendication 12, dans lequel la séquence consensus d'acides aminés de la protéine semblable à Hu-IFN- α contient Arg-Arg-Ala-Ser-Xaa, où Xaa est choisi dans le groupe de Gln, Val ou Leu.
22. Procédé selon la revendication 12, dans lequel la protéine est la protéine semblable à Hu-IFN- α et la protéinekinase est la protéinekinase dépendante de l'AMPc.

23. Procédé selon la revendication 4, comprenant en outre la purification de la protéine phosphorylable semblable à Hu-IFN- α .
24. Procédé selon la revendication 4, comprenant en outre la phosphorylation de la protéine semblable à Hu-IFN- α avec un marqueur radioactif.
25. Procédé selon la revendication 24, dans lequel le marqueur est un atome émettant des rayons bêta.
26. Procédé selon la revendication 25, dans lequel l'atome est choisi dans le groupe constitué du phosphore et du soufre.
27. Procédé selon la revendication 26, dans lequel le marqueur radioactif est ^{32}P ou ^{33}P .
28. Procédé de préparation d'une protéine semblable à Hu-IFN- α ou d'une protéine semblable à Hu-IFN- β , la protéine phosphorylée étant définie dans l'une quelconque des revendications 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21 ou 23, le procédé comprenant l'étape de (a) introduction, dans une protéine qui n'est pas phosphorylable, d'une séquence d'acides aminés reconnaissable par une protéinekinase pour la phosphorylation ou de (b) synthèse de ladite protéine phosphorylable à partir de précurseurs d'acides aminés.
29. Procédé de préparation de la protéine phosphorylable semblable à Hu-IFN- α ayant la séquence d'acides aminés présentée à la figure 3 ou l'une des séquences d'acides aminés identifiées comme $\alpha\text{A-p1}$, $\alpha\text{A-P2}$ et $\alpha\text{A-P3}$, présentées à la figure 8, le procédé comprenant l'étape de (a) introduction, dans une protéine qui n'est pas phosphorylable, d'une séquence d'acides aminés reconnaissable par une protéinekinase pour la phosphorylation ou de (b) synthèse de ladite protéine phosphorylable à partir de précurseurs d'acides aminés.
30. Procédé de préparation de la protéine phosphorylable semblable à Hu-IFN- β de la protéine phosphorylée définie dans la revendication 6, le procédé comprenant l'étape de (a) introduction, dans une protéine qui n'est pas phosphorylable, d'une séquence d'acides aminés reconnaissable par une protéinekinase pour la phosphorylation ou de (b) synthèse de ladite protéine phosphorylable à partir de précurseurs d'acides aminés.
31. Séquence d'ADN recombinant qui code pour une protéine modifiée telle que définie dans la revendication 1.
32. Séquence d'ADN recombinant selon la revendication 31, qui contient une séquence d'ADN qui code pour une protéine fonctionnelle qui normalement n'est pas phosphorylable et une séquence d'ADN qui code pour la séquence consensus d'acides aminés qui est reconnaissable par une protéinekinase et qui contient le site de marquage supposé.
33. Séquence d'ADN recombinant selon la revendication 32, dans laquelle la protéine fonctionnelle est une protéine semblable à un interféron choisie dans le groupe constitué de Hu-IFN- α ou de Hu-IFN- β .
34. Séquence d'ADN recombinant selon la revendication 32, dans laquelle la seconde séquence de nucléotides a un codon qui est commun avec la première séquence de nucléotides.
35. Séquence d'ADN recombinant, qui est choisie dans le groupe constitué des séquences d'ADN codant pour une protéine choisie dans le groupe constitué d'une protéine semblable à Hu-IFN- α ou d'une protéine semblable à Hu-IFN- β , cette protéine ayant été modifiée pour contenir une séquence consensus d'acides aminés reconnaissable par une protéinekinase, comprenant un site phosphorylable supposé.
36. Séquence d'ADN recombinant selon la revendication 35, qui code pour une protéine phosphorylable semblable à Hu-IFN- α .
37. Vecteur de clonage d'ADN recombinant utile pour exprimer une protéine fonctionnelle exogène, ce vecteur incluant la séquence d'ADN qui code pour une protéine semblable à Hu-IFN- α ou une protéine semblable à Hu-IFN- β , cette protéine ayant été modifiée pour contenir une séquence consensus d'acides aminés reconnaissable par une protéinekinase, comprenant un site de phosphorylation supposé, la protéine phosphorylable étant une protéine Hu-IFN- α semblable à un interféron ou une protéine Hu-IFN- β semblable à un interféron.
38. Vecteur de clonage selon la revendication 37, dans lequel la protéine codée est la protéine Hu-IFN- α semblable à un interféron.

39. Vecteur de clonage selon la revendication 37, dans lequel la protéine codée est la protéine Hu-IFN- β semblable à un interféron.
- 5 40. Vecteur de clonage selon la revendication 37, qui contient la séquence d'ADN codant pour une protéine fusionnée Hu-IFN- α /gamma.
41. Vecteur de clonage selon la revendication 37, qui contient la séquence d'ADN codant pour les interférons modifiés dénommés -P1, -P2 et -P3 dans la figure 8, respectivement.
- 10 42. Cellule hôte transformée capable d'exprimer une protéine fonctionnelle, cette cellule hôte contenant un vecteur d'ADN recombinant codant pour une protéine semblable à Hu-IFN- α ou une protéine semblable à Hu-IFN- β qui a été modifiée pour contenir une séquence consensus d'acides aminés reconnaissable par une protéinekinase.
- 15 43. Procédé de fabrication d'une composition biologiquement active, comprenant les étapes de mélange d'un véhicule biologiquement compatible et, d'une quantité biologique efficace, d'une protéine phosphorylée telle que définie dans l'une quelconque des revendications 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26 ou 27.
- 20 44. Procédé selon la revendication 1, dans lequel la protéine phosphorylable possède une propriété biologique de la protéine non phosphorylable.
45. Procédé selon la revendication 1, dans lequel la protéine phosphorylable contient au moins deux segments d'acides aminés, un premier segment qui est la séquence d'acides aminés de la protéine non phosphorylable et un second segment d'acides aminés qui contient la séquence d'acides aminés reconnaissable par la protéinekinase, ce second
25 segment pouvant être incorporé dans le premier segment d'acides aminés ou lié à celui-ci.
46. Procédé selon la revendication 1, dans lequel la protéine phosphorylable est phosphorylée et contient un phosphate stable dans un sérum.
- 30 47. Procédé selon la revendication 1, dans lequel la protéine est un anticorps monoclonal phosphorylable.
48. Procédé selon la revendication 47, dans lequel l'anticorps monoclonal est un anticorps monoclonal chimérique, hybride ou modifié.
- 35 49. Procédé selon la revendication 47, dans lequel l'anticorps monoclonal est spécifique à une tumeur.
50. Procédé selon la revendication 47, dans lequel le site pour la phosphorylation est dans le fragment Fab ou Fab¹ de celui-ci.
- 40 51. Procédé selon l'une quelconque des revendications 47 à 50, comprenant en outre la phosphorylation de l'anticorps monoclonal.
52. Procédé selon l'une quelconque des revendications 47 à 51, comprenant en outre le radiomarquage de l'anticorps monoclonal.
- 45 53. Procédé selon la revendication 1, dans lequel la protéine est une cytokine.
54. Procédé selon la revendication 1, dans lequel la protéine est le facteur de nécrose des tumeurs (TNF) ou l'interleu-
kine-3 (IL-3).
- 50 55. Procédé selon la revendication 54, dans lequel la protéine est choisie dans le groupe constitué du TNF- α et du TNF- β .
- 55 56. Procédé pour préparer une protéine phosphorylable à partir d'une protéine par ailleurs non phosphorylable, comprenant les étapes de :

identification d'une séquence d'acides aminés reconnaissable par une protéinekinase (ci-après "PK") ;
définition et construction d'une séquence de nucléotides qui code pour ladite séquence d'acides aminés recon-

naissable par une PK ;
 détermination d'une séquence de nucléotides codant pour ladite protéine non phosphorylable ;
 construction d'une séquence de nucléotides modifiée codant pour ladite protéine phosphorylable ;
 expression de ladite séquence de nucléotides modifiée pour produire ladite protéine phosphorylable.

- 5
57. Procédé selon la revendication 56, dans lequel ladite étape de construction de ladite séquence de nucléotides modifiée inclut la fixation de ladite séquence de nucléotides codant pour ladite séquence d'acides aminés reconnaissable par une PK à ladite séquence de nucléotides codant pour ladite protéine non phosphorylable.
- 10
58. Procédé selon la revendication 56, dans lequel ladite étape de construction de ladite séquence de nucléotides modifiée inclut l'introduction de ladite séquence de nucléotides codant pour ladite séquence d'acides aminés reconnaissable par une PK dans ladite séquence de nucléotides codant pour ladite protéine non phosphorylable.
- 15
59. Procédé selon la revendication 56, dans lequel ladite séquence d'acides aminés reconnaissable par une PK, identifiée dans ladite étape d'identification, contient Ser.
60. Procédé selon la revendication 56, dans lequel ladite séquence d'acides aminés reconnaissable par une PK, identifiée dans ladite étape d'identification, contient un acide aminé choisi dans le groupe constitué de Ser, Thr ou Tyr.
- 20
61. Procédé selon la revendication 56, dans lequel ladite séquence d'acides aminés reconnaissable par une PK identifiée dans ladite étape d'identification est Arg-Arg-Ala-Ser-.
- 25
62. Procédé selon la revendication 56, dans lequel ladite séquence d'acides aminés reconnaissable par une PK identifiée dans ladite étape d'identification contient Arg-Arg-Ala-Ser-Xaa, où Xaa est choisi dans le groupe de tous les acides aminés.
63. Procédé selon la revendication 56, dans lequel la protéine kinase de ladite étape d'identification est la protéine kinase dépendant de l'AMPc.
- 30
64. Procédé selon la revendication 56, dans lequel ladite protéine phosphorylable est phosphorylée avec un marqueur radioactif.

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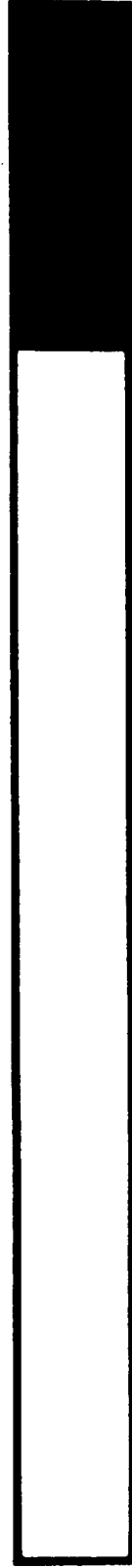
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HYBRID Hu-IFN-αA/δ

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Hu-IFN-αA
(165 AMINO ACIDS)

Hu-IFN-δ
(16 AMINO ACIDS)

FIG.1

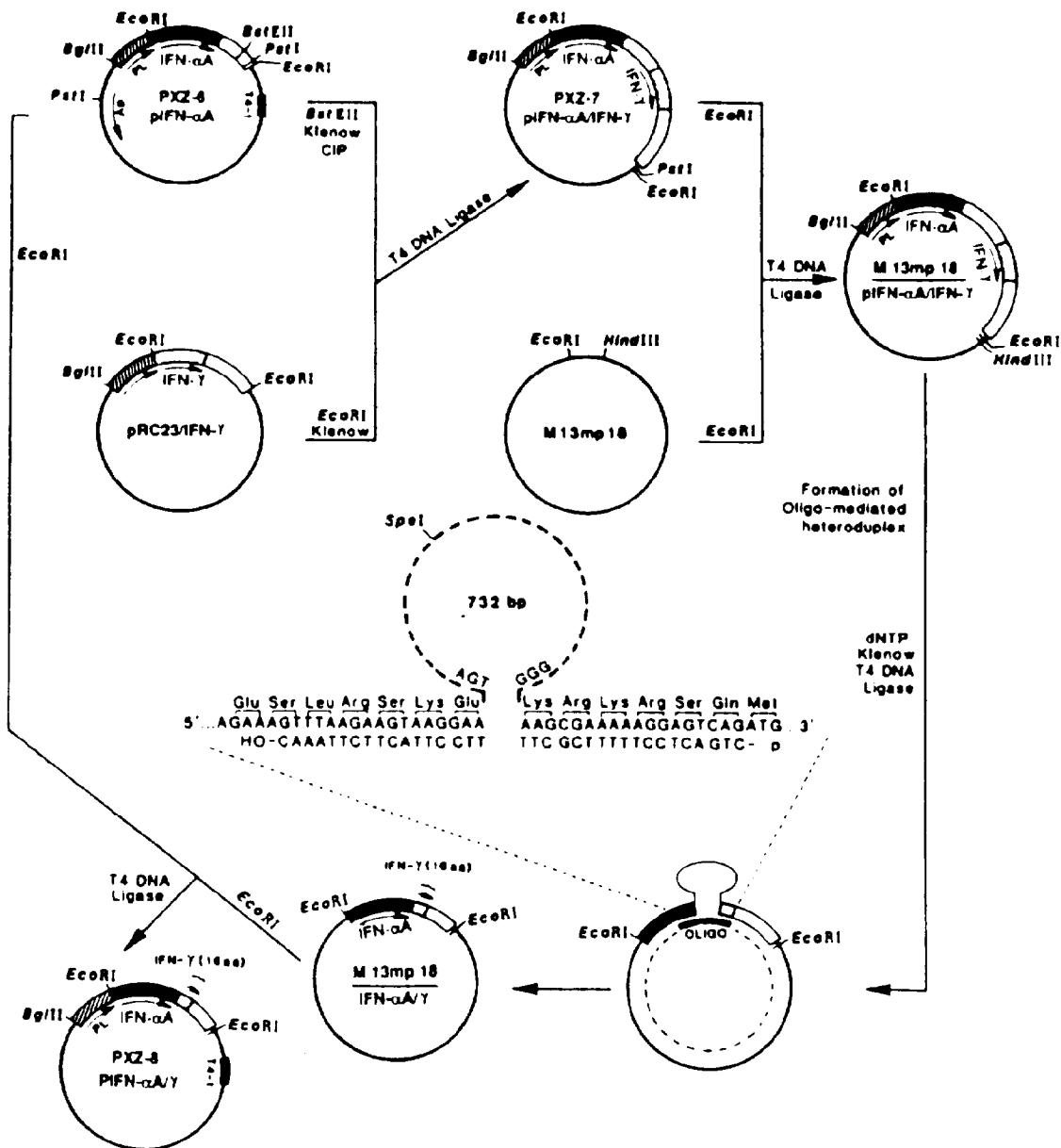
CONSTRUCTION OF EXPRESSION VECTOR FOR Hu-IFN- α A/ γ HYBRID

FIG.2

NUCLEOTIDE AND AMINO SEQUENCES OF HYBRID HU-IFN- α /8

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5' 1 20
   MET CYS ASP LEU PRO GLN THR HIS SER LEU GLY SER ARG ARG THR LEU MET LEU LEU ALA GLN MET ARG LYS ILE
   ATG TGT GAT CTG CCT CAA ACC CAC AGC CTG GGT AGC AGG AGG ACC TTG ATG CTC CTG GCA CAG ATG AGG AAA ATC

   30 40
   SER LEU PHE SER CYS LEU LYS ASP ARG HIS ASP PHE GLY PHE PRO GLN GLU PHE GLY ASN GLN PHE GLN LYS
   TCT CTT TTC TCC TGC TTG AAG GAC AGA CAT GAC TTT GGA TTT CCC CAG GAG GAG TTT GGC AAC CAG TTC CAA AAG

   50 60 70
   ALA GLU THR ILE PRO VAL LEU HIS GLU MET ILE GLN GLN ILE PHE ASN LEU PHE SER THR LYS ASP SER SER ALA
   GCT GAA ACC ATG CCT GTC CTC CAT GAG ATG ATC CAG CAG ATC TTC AAT CTC TTC AGC ACA AAG GAC TCA TCT GCT

   80 90 100 110 120
   ALA THR ASP GLU THR LEU LEU ASP LYS PHE TYR THR GLU LEU TYR GLN GLN LEU ASN ASP LEU GLU ALA CYS VAL
   GCT TGG GAT GAG ACC CTC CTA GAC AAA TTC TAC ACT GAA CTC TAC CAG CAG CTG AAT GAC CTG GAA GCC TGT GTG

   130 140 150 160 170
   ILE GLN GLY VAL GLY VAL THR GLU THR PRO LEU MET LYS GLU ASP SER ILE LEU ALA VAL ARG LYS TYR PHE GLN
   ATA CAG GGG GTG GGG GTG ACA GAG ACT CCC CTG ATG AAG GAG GAC TCC ATT CTG GCT GTG AGG AAA TAC TTC CAA

   180 190 200
   ARG ILE THR LEU TYR LEU LYS GLU LYS LYS TYR SER PRO CYS ALA TRP GLU VAL VAL ARG ALA GLU ILE MET ARG
   AGA ATC ACT CTC TAT CTG AAA GAG AAG AAA TAC AGC CCT TGT GCC TGG GAG GTT GTC AGA GCA GAA ATC ATG AGA

   210 220 230 240 250
   SER PHE SER LEU SER THR ASN LEU GLN GLU SER LEU ARG SER LYS GLU LYS ARG LYS ARG SER GLN MET LEU PHE
   TCT TTT TCT TGT TCA ACA AAC AAG TTG CAA GAA AGT TTA AGA AGT AAG GAA AAG CGA AAA AGG AGT CAG ATG CTG TTT

   260 270 280 290 300
   GLN GLY ARG ARG ALA SER GLN END
   CAA GGT CGA AGA GCA TCC CAG TAA TGGTGGCTGCTGCAATATTG 3'

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FIG.3

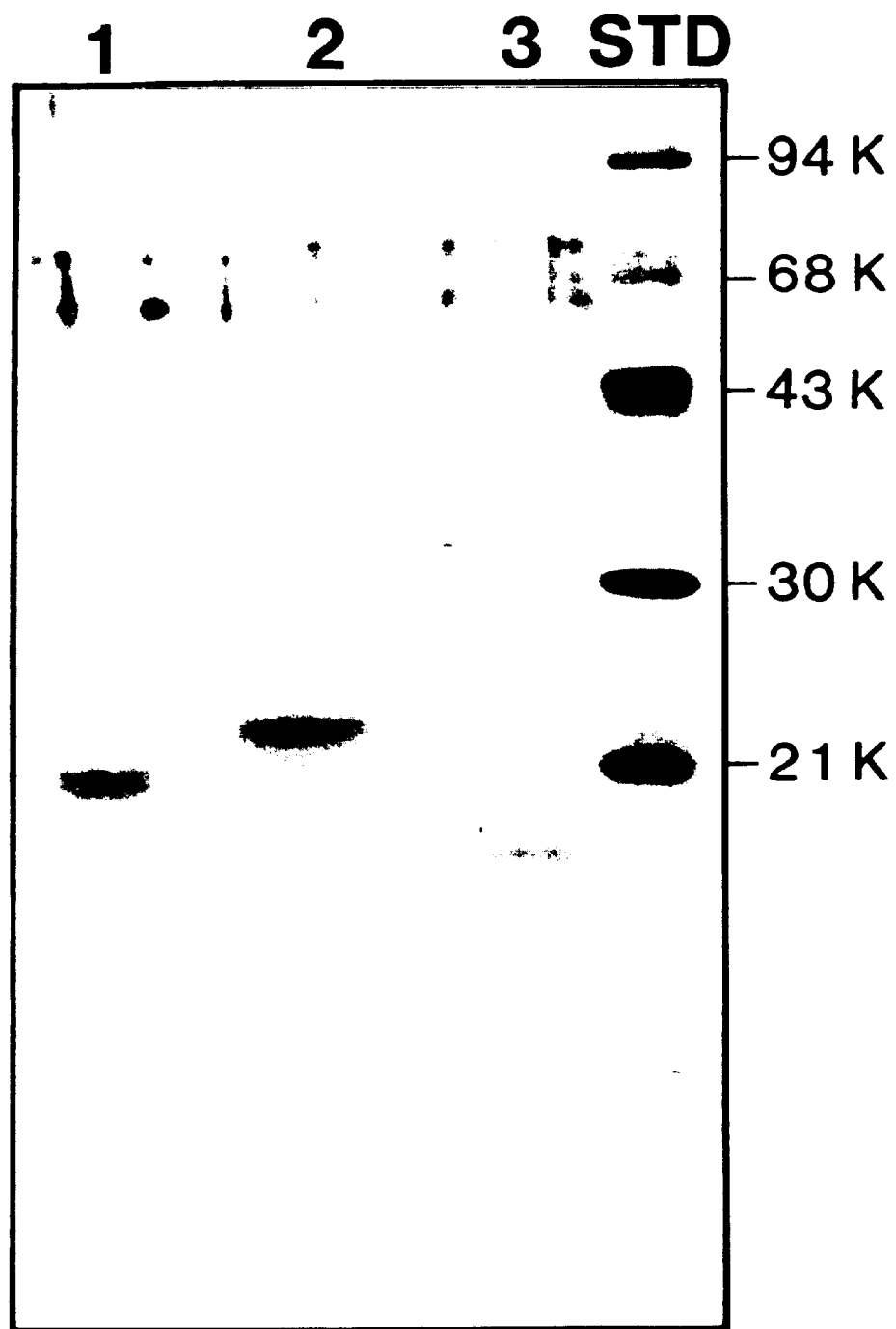
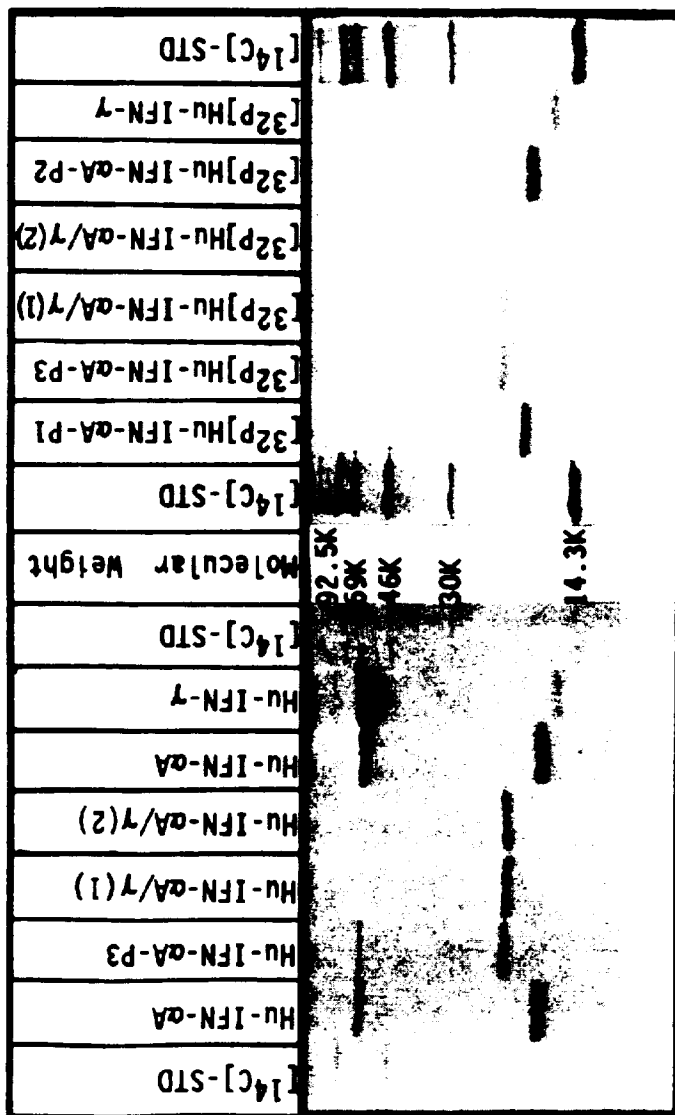


FIG. 4

FIG.5



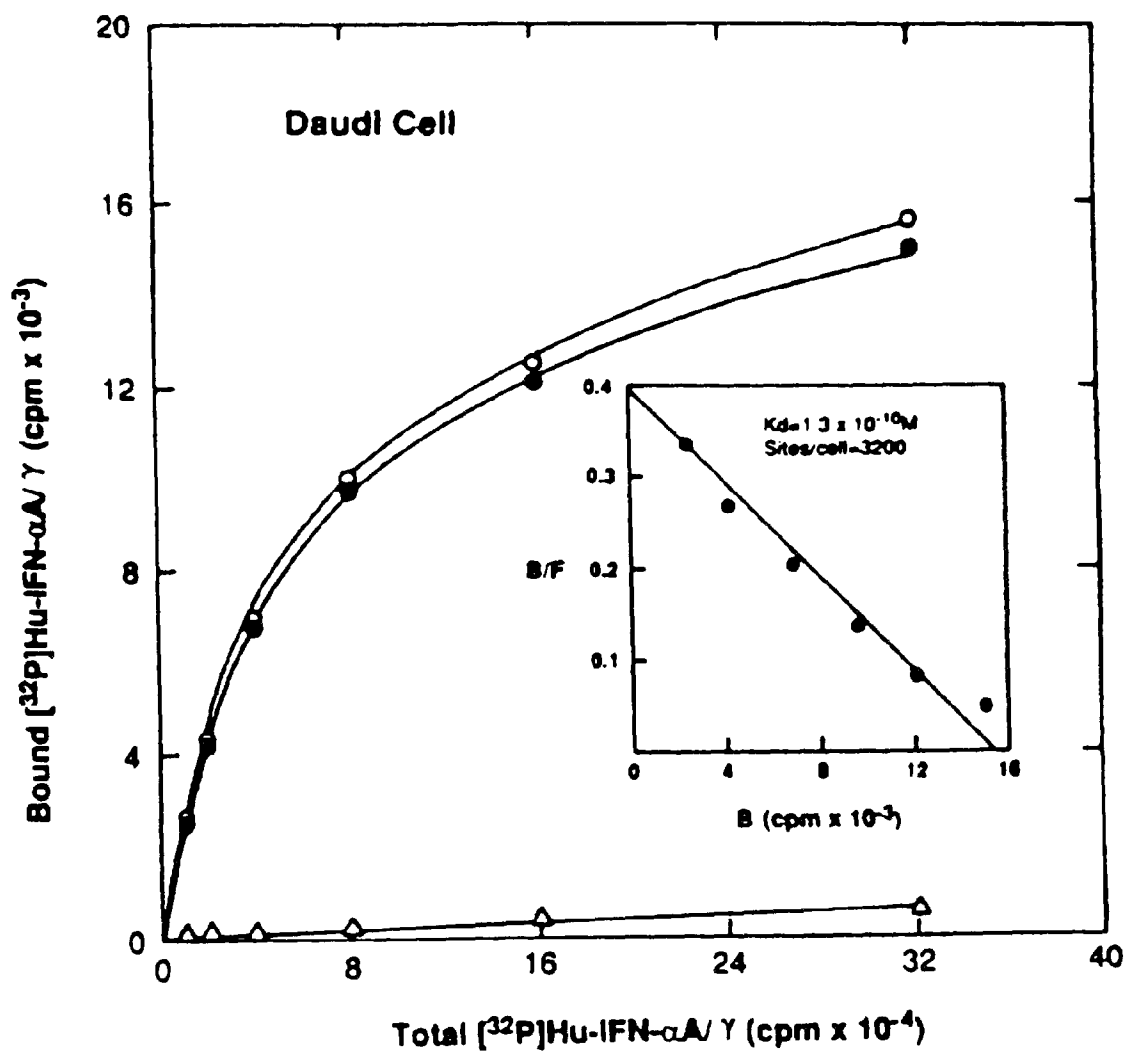


FIG. 6

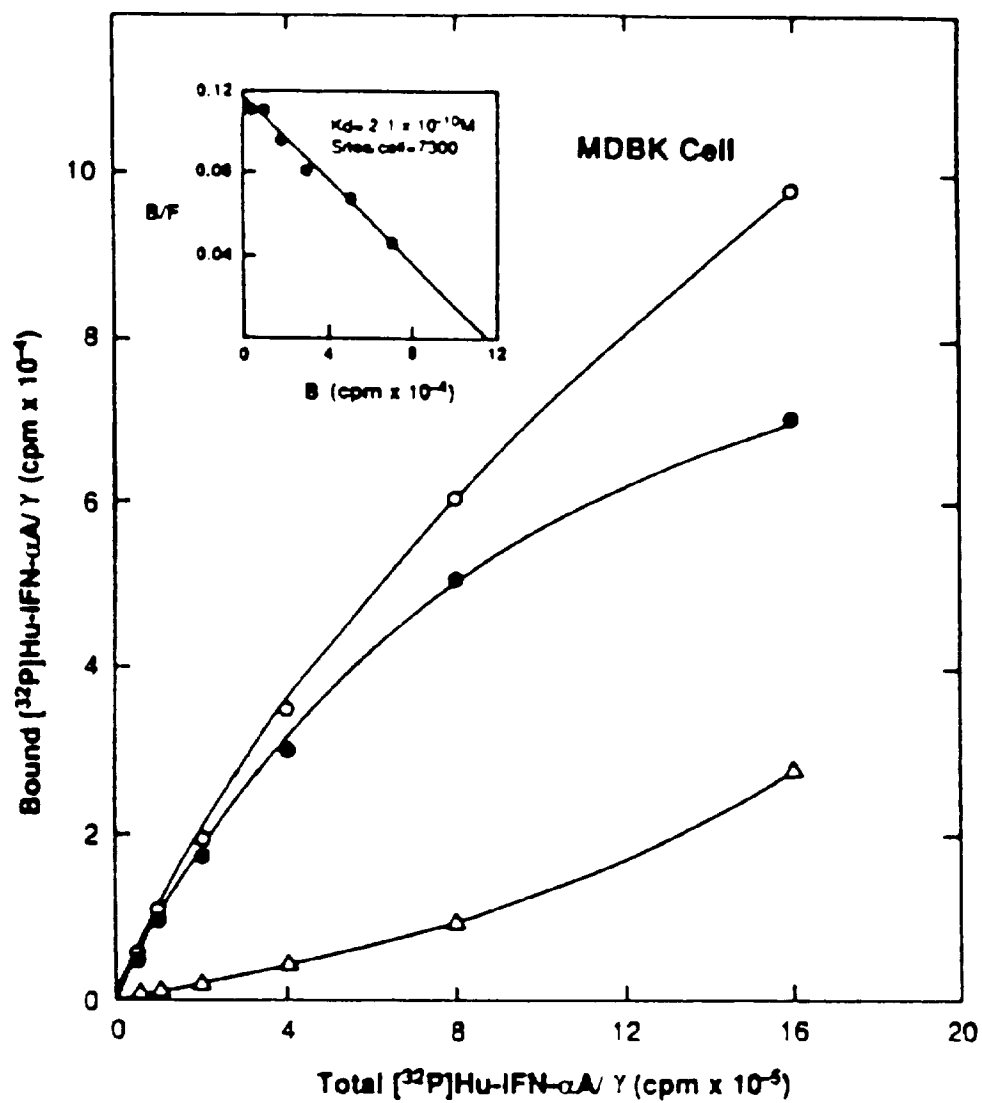


FIG. 7

COMPARISON OF NUCLEOTIDE AND AMINO ACID SEQUENCES OF
Hu-IFN- α A AND ANALOGUES WITH PHOSPHORYLATION SITES

α A	155	5'...	ACA AAC	160	TTG CAA	165	AGT TTA	AGA AGT	170	AAG GAA	TGA
		Thr	Asn	Leu	Gln	Glu	Ser	Leu	Arg	Ser	Lys
											Glu
											END
α A-P1	155	5'...	ACA AAC	160	TTG CAA	165	AGT TTA	AGA AGT	170	AGG GCA	AGT GTT
		Thr	Asn	Leu	Gln	Glu	Ser	Leu	Arg	Ser	Lys
											Ala
											Ala
											END
α A-P2	155	5'...	ACA AAC	160	TTG CAA	165	AGT TTA	AGA AGG	167	GCA AGT	TTA GCA
		Thr	Asn	Leu	Gln	Arg	Ser	Leu	Arg	Ala	Ser
											Leu
											Ala
											END
α A-P3	155	5'...	ACA AAC	160	TTG CAA	165	AGT TTA	AGA AGT	175	AGA AGT	AGA AGG
		Thr	Asn	Leu	Gln	Glu	Ser	Leu	Arg	Ser	Arg
											Glu
											Ser
											Leu
											Arg
											Ala
											Ser
											Val
											Ala
											END

FIG.8

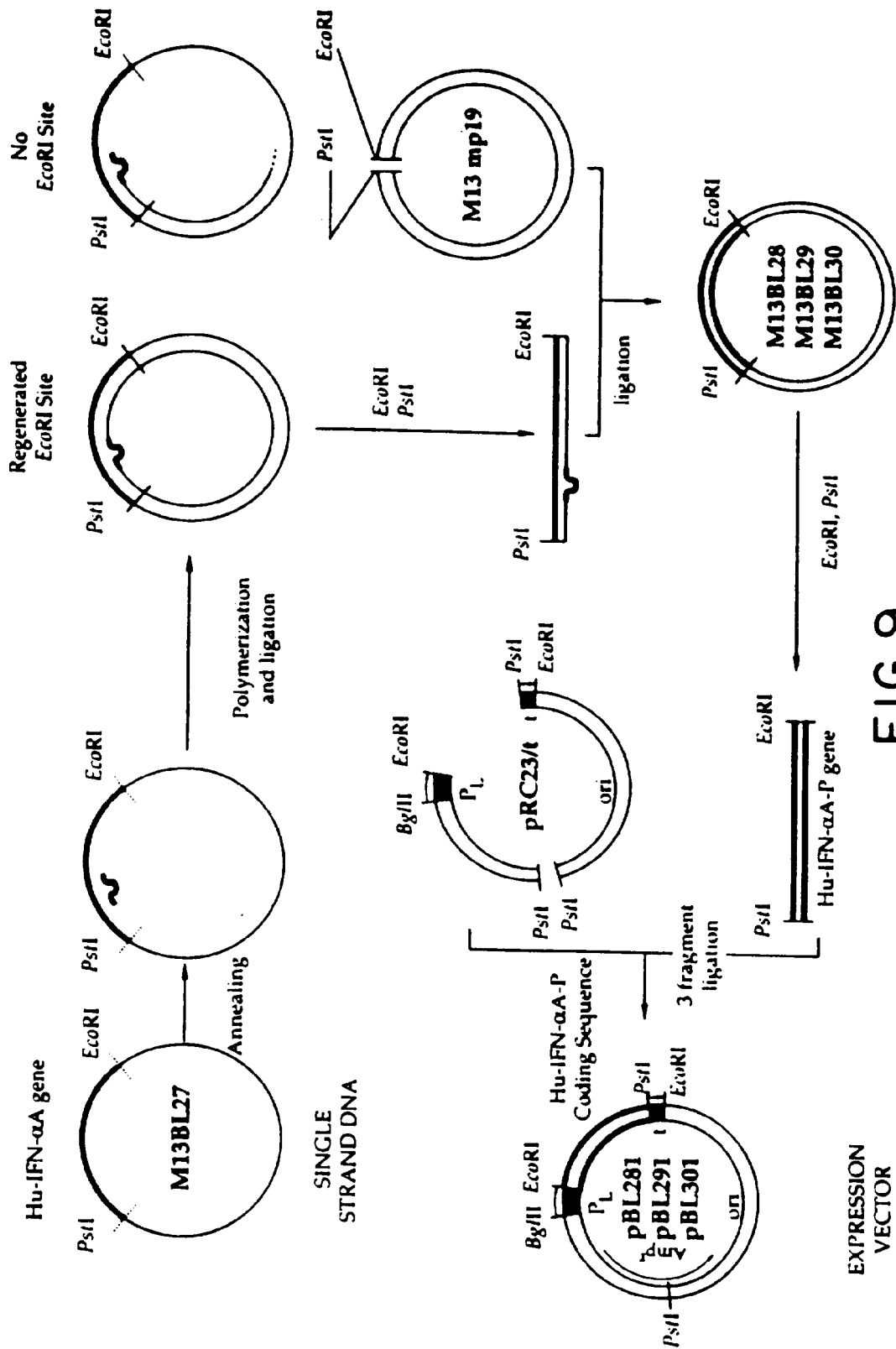


FIG.9

FIG.10

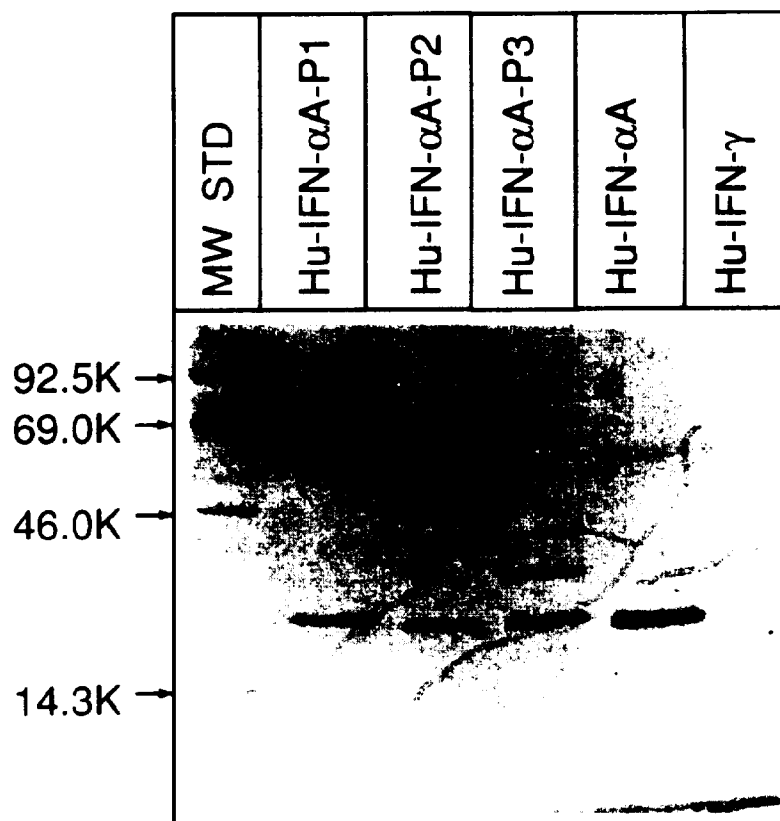
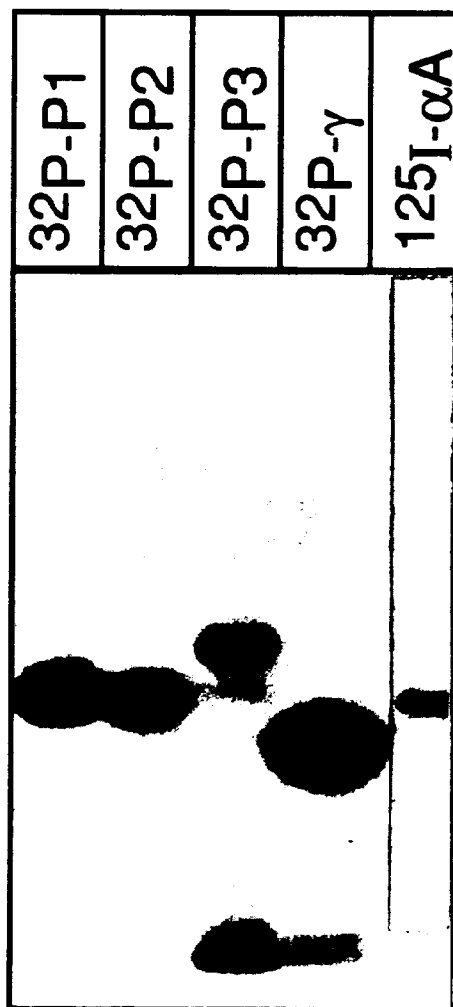
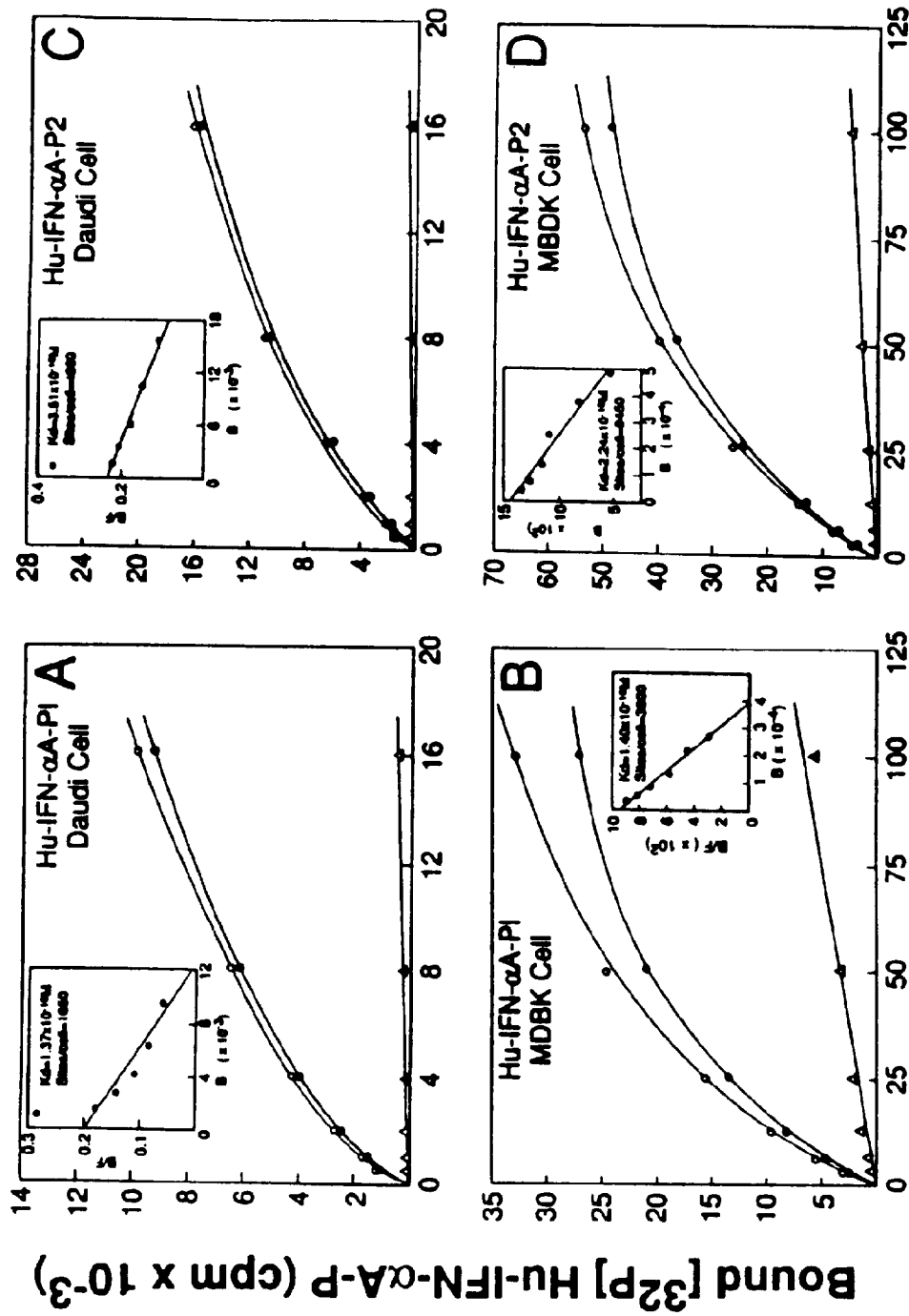


FIG. 11





$[^{32}\text{P}]$ Hu-IFN- α A-P Concentration ($\text{cpm} \times 10^{-4}$)

FIG.12

FIG.13

